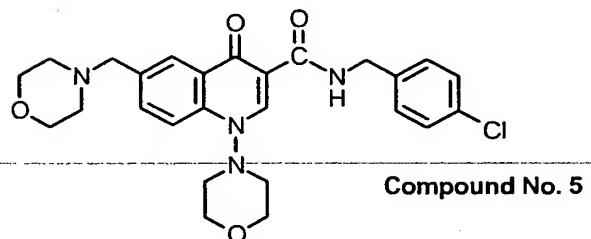
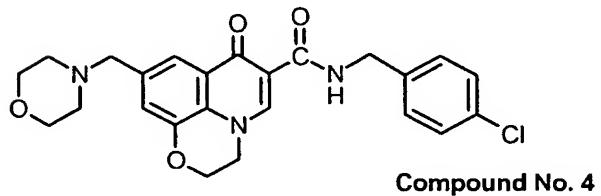
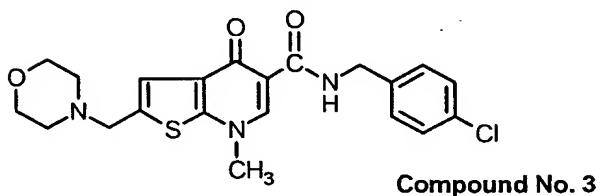
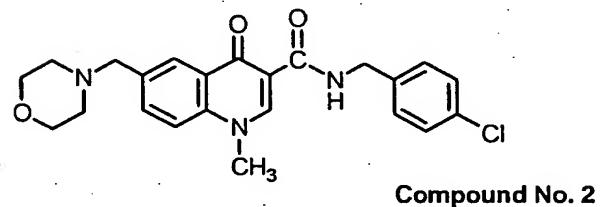
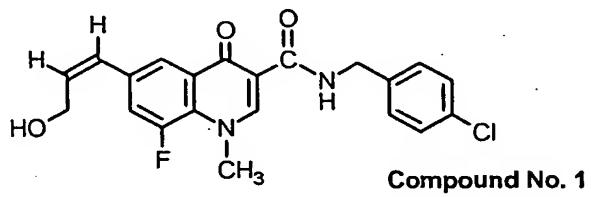
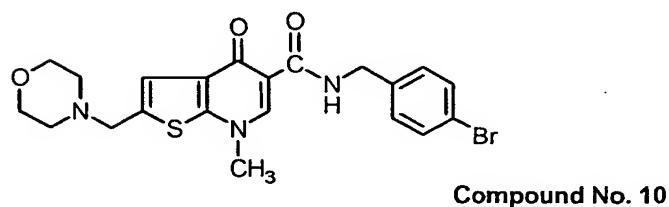
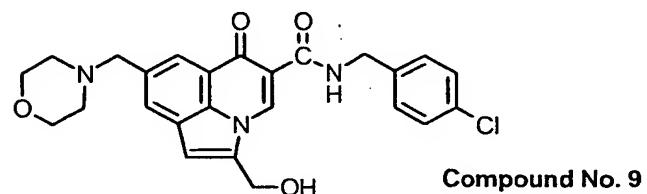
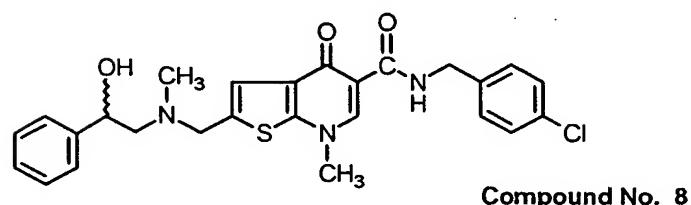
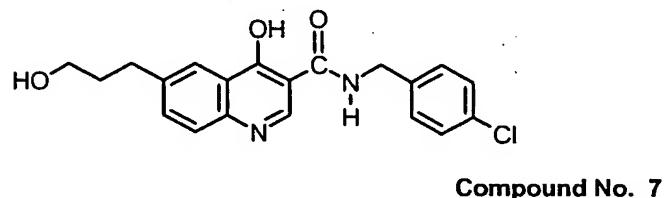
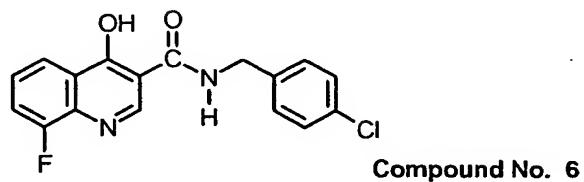


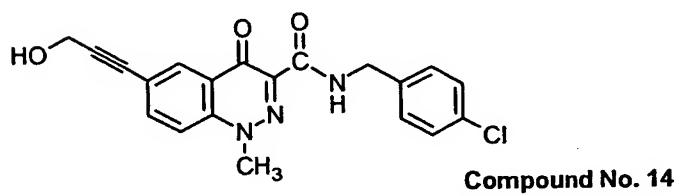
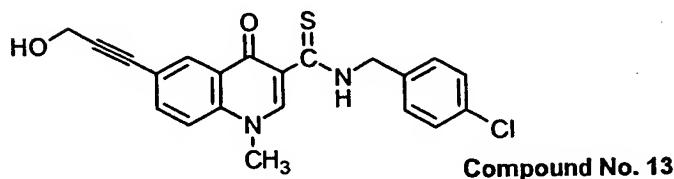
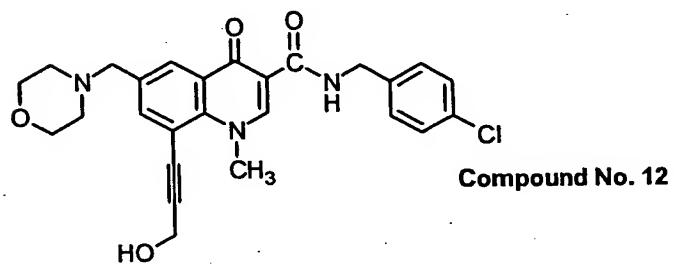
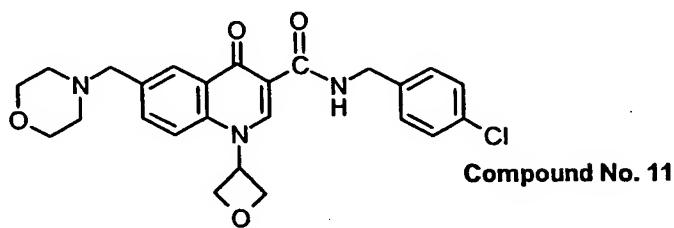
5 **Figure 1A 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds**



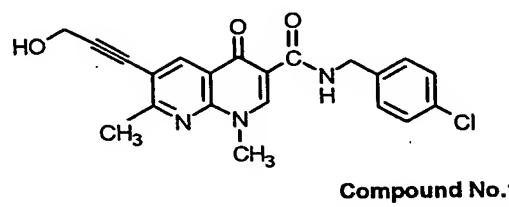
5 **Figure 1B 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds**



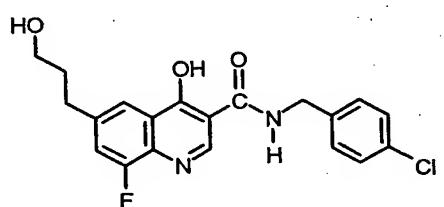
5 **Figure 1C 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds**



5 **Figure 1D 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds**



Compound No.15



Compound 17

Figure 4A Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Aligned by Amino Acid Homology*

	HSV2-MS	MFCAGGPTS PGGKSAARAA SGFFAPHNPR GATQTAPPPC RRQNFTYNPHL	-50
	HSV2-186	MFCAGGPPAS PGGKSAARAA SGFFAPHNPR GATQTAPPPC RRQNFTYNPHL	-50
5	HSV1-Kos	MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGR.GPPPC LRQNFTYNPYL	-49
	HSV1-Patton	MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGR.GPPPC LRQNFTYNPYL	-49
	HSV1-DJL	MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGR.GPPPC LRQNFTYNPYL	-49
	HSV1-F	MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGR.GPPPC LRQNFTYNPYL	-49
10	HSV2-MS	AQTGTQPKAP GPAQRHTYYS ECDEFRFIAPI RSLDEDAPAE QRTGVHDGRL	-100
	HSV2-186	AQTGTQPKAP GPAQRHTYYS ECDEFRFIAPI RSLDEDAPAE QRTGVHDGRL	-100
	HSV1-Kos	APVGTQQKPT GPTQRHTYYS ECDEFRFIAPI RVLDEDAPPE KAGVHDGHL	-99
	HSV1-Patton	APVGTQQKPT GPTQRHTYYS ECDEFRFIAPI RVLDEDAPPE KAGVHDGHL	-99
	HSV1-DJL	APVGTQQKPT GPTQRHTYYS ECDEFRFIAPI RVLDEDAPPE KAGVHDGHL	-99
15	HSV1-F	APVGTQQKPT GPTQRHTYYS ECDEFRFIAPI RVLDEDAPPE KAGVHDGHL	-99
	HSV2-MS	RRAPKVCAGG DERDVLRVGP EGFWPRLRL WGGADHAPKG FDPTVTVFHV	-150
	HSV2-186	RRAPKVCAGG DERDVLRVGP EGFWPRLRL WGGADHAPEG FDPTVTVFHV	-150
	HSV1-Kos	KRAPKVCAGG DERDVLRVGS GGFWRSSRL WGGVDHAPAG FNPTVTVFHV	-149
20	HSV1-Patton	KRAPKVCAGG DERDVLRVGS GGFWRSSRL WGGVDHAPAG FNPTVTVFHV	-149
	HSV1-DJL	KRAPKVCAGG DERDVLRVGS GGFWRSSRL WGGVDHAPAG FNPTVTVFHV	-149
	HSV1-F	KRAPKVCAGG DERDVLRVGS GGFWRSSRL WGGVDHAPAG FNPTVTVFHV	-149
	HSV2-MS	YDILEHVEHA YSMRAAQLHE RFMDAITPAG TVITLLGLTP EGHRAVHVY	-200
25	HSV2-186	YDILEHVEHA YSMRAAQLHE RFMDAITPAG TVITLLGLTP EGHRAVHVY	-200
	HSV1-Kos	YDILENVEHA YGMRAAQFHA RFMDAITPTG TVITLLGLTP EGHRAVHVY	-199
	HSV1-Patton	YDILENVEHA YGMRAAQFHA RFMDAITPTG TVITLLGLTP EGHRAVHVY	-199
	HSV1-DJL	YDILENVEHA YGMRAAQFHA RFMDAITPTG TVITLLGLTP EGHRAVHVY	-199
	HSV1-F	YDILENVEHA YGMRAAQFHA RFMDAITPTG TVITLLGLTP EGHRAVHVY	-199
30	HSV2-MS	GTRQYFYMNK AEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE	-250
	HSV2-186	GTRQYFYMNK AEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE	-250
	HSV1-Kos	GTRQYFYMNK EEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE	-249
	HSV1-Patton	GTRQYFYMNK EEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE	-249
35	HSV1-DJL	GTRQYFYMNK EEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE	-249
	HSV1-F	GTRQYFYMNK EEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE	-249
	HSV2-MS	AEVVERADVY YYETRPTLYY RVFVRSGRAL AYLCDNFCPA IRKYEGGVDA	-300
	HSV2-186	AEVVERADVY YYETRPTLYY RVFVRSGRAL AYLCDNFCPA IRKYEGGVDA	-300
40	HSV1-Kos	AEVVERTDVY YYETRPAFLY RVYVRSGRVL SYLCDNFCPA IKKYEGGVDA	-299
	HSV1-Patton	AEVVERTDVY YYETRPAFLY RVYVRSGRVL SYLCDNFCPA IKKYEGGVDA	-299
	HSV1-DJL	AEVVERTDVY YYETRPAFLY RVYVRSGRVL SYLCDNFCPA IKKYEGGVDA	-299
	HSV1-F	AEVVERTDVY YYETRPAFLY RVYVRSGRVL SYLCDNFCPA IKKYEGGVDA	-299
45	HSV2-MS	TTRFILDNP G FVTFGWYRLK PGRGNAPAQP RPPTAFTGSS DVEFNCTADN	-350
	HSV2-186	TTRFILDNP G FVTFGWYRLK PGRGNAPAQP RPPTAFTGSS DVEFNCTADN	-350
	HSV1-Kos	TTRFILDNP G FVTFGWYRLK PGRNNTLAQP RAPMAFGTSS DVEFNCTADN	-349
	HSV1-Patton	TTRFILDNP G FVTFGWYRLK PGRNNTLAQP RAPMAFGTSS DVEFNCTADN	-349
	HSV1-DJL	TTRFILDNP G FVTFGWYRLK PGRNNTLAQP RAPMAFGTSS DVEFNCTADN	-349
50	HSV1-F	TTRFILDNP G FVTFGWYRLK PGRNNTLAQP RAPMAFGTSS DVEFNCTADN	-349
	HSV2-MS	LAVEGAMCDL PAYKLMCFDI ECKAGGEDEL AFPVAERPED LVIQISCLLY	-400
	HSV2-186	LAVEGAMCDL PAYKLMCFDI ECKAGGEDEL AFPVAERPED LVIQISCLLY	-400
	HSV1-Kos	LAIEGGMSDL PAYKLMCFDI ECKAGGEDEL AFPVAGHPED LVIQISCLLY	-399
55	HSV1-Patton	LAIEGGMSDL PAYKLMCFDI ECKAGGEDEL AFPVAGHPED LVIQISCLLY	-399
	HSV1-DJL	LAIEGGMSDL PAYKLMCFDI ECKAGGEDEL AFPVAGHPED LVIQISCLLY	-399
	HSV1-F	LAIEGGMSDL PAYKLMCFDI ECKAGGEDEL AFPVAGHPED LVIQISCLLY	-399
	HSV2-MS	DLSTTALEHI LLFSLGSCDL PESHLSDLAS RGLPAPVVLE FDSEFEMLLA	-450

Figure 4B Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Aligned by Amino Acid Homology*

	HSV2-186	DLSTTALLEHI LLFSLGSCDL PESHLSDLAS RGLPAPVVLE FDSEFEMLLA -450
5	HSV-Kos	DLSTTALLEHV LLFSLGSCDL PESHLNELAA RGLPTPVVLE FDSEFEMLLA -449
	HSV1-Patton	DLSTTALLEHV LLFSLGSCDL PESHLNELAA RGLPTPVVLE FDSEFEMLLA -449
	HSV1-DJL	DLSTTALLEHV LLFSLGSCDL PESHLNELAA RGLPTPVVLE FDSEFEMLLA -449
	HSV1-F	DLSTTALLEHV LLFSLGSCDL PESHLNELAA RGLPTPVVLE FDSEFEMLLA -449
10	HSV2-MS	FMTFVKQYGP EFVTGYNIIN FDWPFLVTLKL TEIYKVPLDG YGRMNGRGVF -500
	HSV2-186	FMTFVKQYGP EFVTGYNIIN FDWPFLVTLKL TEIYKVPLDG YGRMNGRGVF -500
	HSV-Kos	FMTLVKQYGP EFVTGYNIIN FDWPFLLAKL TDIYKVPLDG YGRMNGRGVF -499
	HSV1-Patton	FMTLVKQYGP EFVTGYNIIN FDWPFLLAKL TDIYKVPLDG YGRMNGRGVF -499
	HSV1-DJL	FMTLVKQYGP EFVTGYNIIN FDWPFLLAKL TDIYKVPLDG YGRMNGRGVF -499
15	HSV1-F	FMTLVKQYGP EFVTGYNIIN FDWPFLLAKL TDIYKVPLDG YGRMNGRGVF -499
	HSV2-MS	RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKVKLSSYK LNAVAEAVLK -550
	HSV2-186	RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKVKLSSYK LNAVAEAVLK -550
	HSV-Kos	RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKIKLSSYK LNAVAEAVLK -549
20	HSV1-Patton	RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKIKLSSYK LNAVAEAVLK -549
	HSV1-DJL	RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKIKLSSYK LNAVAEAVLK -549
	HSV1-F	RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKIKLSSYK LNAVAEAVLK -549
25	HSV2-MS	DKKKDLSYRD IPAYYAS GPA QRGVIGEYCV QDSLLVGQLF FKFLPHLELS -600
	HSV2-186	DKKKDLSYRD IPAYYAS GPA QRGVIGEYCV QDSLLVGQLF FKFLPHLELS -600
	HSV-Kos	DKKKDLSYRD IPAYYAAGPA QRGVIGEYCI QDSLLVGQLF FKFLPHLELS -599
	HSV1-Patton	DKKKDLSYRD IPAYYAAGPA QRGVIGEYCI QDSLLVGQLF FKFLPHLELS -599
	HSV1-DJL	DKKKDLSYRD IPTYYAAGPA QRGVIGEYCI QDSLLVGQLF FKFLPHLELS -599
	HSV1-F	DKKKDLSYRD IPAYYAAGPA QRGVIGEYCI QDSLLVGQLF FKFLPHLELS -599
30	HSV2-MS	AVARLAGINI TRTIYDGQQI RVFTCLLR LA GQKGFI LPDT QGRFRGLDKE -650
	HSV2-186	AVARLAGINI TRTIYDGQQI RVFTCLLR LA GQKGFI LPDT QGRFRGLDKE -650
	HSV-Kos	AVARLAGINI TRTIYDGQQI RVFTCLLR LA DQKGFI LPDT QGRFRGAGGE -649
	HSV1-Patton	AVARLAGINI TRTIYDGQQI RVFTCLLR LA DQKGFI LPDT QGRFRGAGGE -649
35	HSV1-DJL	AVARLAGINI TRTIYDGQQI RVFTCLLR LA DQKGFI LPDT QGRFRGAGGE -649
	HSV1-F	AVARLAGINI TRTIYDGQQI RVFTCLLR LA DQKGFI LPDT QGRFRGGGGE -649
	HSV2-MS	APKRPAVPRG EGERPGDGNG DEDKDDDE.. DEDGDERE.E VARETGGRHV -697
	HSV2-186	APKRPAVPRG EGERPGDGNG DEDKDDDEDG DEDGDERE.E VARETGGRHV -697
40	HSV-Kos	APKRPAAAARE DEERP..... EEEGEDEDER EEGGGEREPE GARETAGRHV -694
	HSV1-Patton	APKRPAAAARE DEERP..... EEEGEDEDER EEGGGEREPE GARETAGRHV -694
	HSV1-DJL	APKRPAAAARE DEERP..... EEEGEDENER EEGGGEREPE GARETAGRHV -694
	HSV1-F	APKRPAAAARE DEERP..... EEEGEDEDER EEGGGEREPE GARETAGRHV -694
45	HSV2-MS	GYQGARVLDP TSGFHVPVV VFDFASLYPS IIQAHNLCS TLSLRPEAVA -747
	HSV2-186	GYQGARVLDP TSGFHVPVV VFDFASLYPS IIQAHNLCS TLSLRPEAVA -749
	HSV-Kos	GYQGARVLDP TSGFHVPVV VFDFASLYPS IIQAHNLCS TLSLRADAVA -744
	HSV1-Patton	GYQGARVLDP ISGFHVPVV VFDFASLYPS IIQAHNLCS TLSLRADAVA -744
	HSV1-DJL	GYQGARVLDP TSGFHVPVV VFDFASLYPS IIQAHNLCS TLSLRADAVA -744
50	HSV1-F	GYQGARVLDP TSGFHVPVV VFDFASLYPS IIQAHNLCS TLSLRADAVA -744
	HSV2-MS	HLEADR DYLE IEVGGRLFFF VKAHVRESLL SILLRDWLAM RKQIRSRIPO -797
	HSV2-186	HLEADR DYLE IEVGGRLFFF VKAHVRESLL SILLRDWLAM RKQIRSRIPO -799
	HSV-Kos	HLEAGKDYLE IEVGGRLFFF VKAHVRESLL SILLRDWLAM RKQIRSRIPO -794
55	HSV1-Patton	HLEAGKDYLE IEVGGRLFFF VKAHVRESLL SILLRDWLAM RKQIRSRIPO -794
	HSV1-DJL	HLEAGKDYLE IEVGGRLFFF VKAHVRESLL SILLRDWLAM RKQIRSRIPO -794
	HSV1-F	HLEAGKDYLE IEVGGRLFFF VKAHVRESLL SILLRDWLAM RKQIRSRIPO -794
60	HSV2-MS	STPPEAVLLD KQQAAIKVVC NSVYGFTGVQ HGLLPCLHVA ATVTTIGREM -847
	HSV2-186	SPPEEA VLLD KQQAAIKVVC NSVYGFTGVQ HGLLPCLHVA ATVTTIGREM -849
	HSV-Kos	SSPEEA VLLD KQQAAIKVVC NSVYGFTGVQ HGLLPCLHVA ATVTTIGREM -844

Figure 4C Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Aligned by Amino Acid Homology*

	HSV1-Patton	SSPEEAVLLD KQQAAIKVVC NSVYGFVGQ HGLLPCLHVA ATVTTIGREM -844
5	HSV1-DJL	SSPEEAVLLD KQQAAIKVVC NSVYGFVGQ HGLLPCLHVA ATVTTIGREM -844
	HSV1-F	SSPEEAVLLD KQQAAIKVVC NSVYGFVGQ HGLLPCLHVA ATVTTIGREM -844
	HSV2-MS	LLATRAYVHA RWAEDQQLA DFPEAAGMRA PGPYSMRIIY GDTDSIFVLC -897
10	HSV2-186	LLATRAYVHA RWAEDQQLA DFPEAAGMRA PGPYSMRIIY GDTDSIFVLC -899
	HSV-Kos	LLATREYVHA RWAFAEQLLA DFPEAADMRA PGPYSMRIIY GDTDSIFVLC -894
	HSV1-Patton	LLATREYVHA RWAFAEQLLA DFPEAADMRA PGPYSMRIIY GDTDSIFVLC -894
	HSV1-DJL	LLATREYVHA RWAFAEQLLA DFPEAADMRA PGPYSMRIIY GDTDSIFVLC -894
	HSV1-F	LLATREYVHA RWAFAEQLLA DFPEAADMRA PGPYSMRIIY GDTDSIFVLC -894
15	HSV2-MS	RGLTAAGLVA MGDKMASHIS RALFLPPIKL ECEKTFTKLL LIAKKYIGV -947
	HSV2-186	RGLTAAGLVA MGDKMASHIS RALFLPPIKL ECEKTFTKLL LIAKKYIGV -949
	HSV-Kos	RGLTAAGLTA MGDKMASHIS RALFLPPIKL ECEKTFTKLL LIAKKYIGV -944
	HSV1-Patton	RGLTAAGLTA MGDKMASHIS RALFLPPIKL ECEKTFTKLL LIAKKYIGV -944
	HSV1-DJL	RGLTAAGLTA VGDKMASHIS RALFLPPIKL ECEKTFTKLL LIAKKYIGV -944
20	HSV1-F	RGLTAAGLTA VGDKMASHIS RALFLSPIKL ECEKTFTKLL LIAKKYIGV -944
	HSV2-MS	IYGGKMLIKG VDLVRKNNC FINRTSRALV DLLFYDDTVS GAAAALAERP -997
	HSV2-186	IYGGKMLIKG VDLVRKNNC FINRTSRALV DLLFYDDTVS GAAAALAERP -999
25	HSV-Kos	IYGGKMLIKG VDLVRKNNC FINRTSRALV DLLFYDDTVS GAAAALAERP -994
	HSV1-Patton	IYGGKMLIKG VDLVRKNNC FINRTSRALV DLLFYDDTVS GAAAALAERP -994
	HSV1-DJL	IYGGKMLIKG VDLVRKNNC FINRTSRALV DLLFYDDTVS GAAAALAERP -994
	HSV1-F	IYGGKMLIKG VDLVRKNNC FINRTSRALV DLLFYDDTVS GAAAALAERP -994
	HSV2-MS	AEEWLARPLP EGLQAFGAVL VDAHRRITDP ERDIQDFVLT AELSRRHPRAY -1047
30	HSV2-186	AEEWLARPLP EGLQAFGAVL VDAHRRITDP ERDIQDFVLT AELSRRHPRAY -1049
	HSV-Kos	AEEWLARPLP EGLQAFGAVL VDAHRRITDP ERDIQDFVLT AELSRRHPRAY -1044
	HSV1-Patton	AEEWLARPLP EGLQAFGAVL VDAHRRITDP ERDIQDFVLT AELSRRHPRAY -1044
	HSV1-DJL	AEEWLARPLP EGLQAFGAVL VDAHRRITDP ERDIQDFVLT AELSRRHPRAY -1044
	HSV1-F	AEEWLARPLP EGLQAFGAVL VDAHRRITDP ERDIQDFVLT AELSRRHPRAY -1044
35	HSV2-MS	TNKRLAHLTV YYKLMARRAQ VPSIKDRIPY VIVAQTREVE ETVARLAALR -1097
	HSV2-186	TNKRLAHLTV YYKLMARRAQ VPSIKDRIPY VIVAQTREVE ETVARLAALR -1099
	HSV-Kos	TNKRLAHLTV YYKLMARRAQ VPSIKDRIPY VIVAQTREVE ETVARLAALR -1094
	HSV1-Patton	TNKRLAHLTV YYKLMARRAQ VPSIKDRIPY VIVAQTREVE ETVARLAALR -1094
40	HSV1-DJL	TNKRLAHLTV YYKLMARRAQ VPSIKDRIPY VIVAQTREVE ETVARLAALR -1094
	HSV1-F	TNKRLAHLTV YYKLMARRAQ VPSIKDRIPY VIVAQTREVE ETVARLAALR -1094
	HSV2-MS	ELDAAAAPGDE PAPPAALPSP AKRPRETPSH ADPPGGASKP RKLLVSELAE -1147
	HSV2-186	ELDAAAAPGDE PAPPAALPSP AKRPRETPSH ADPPGGASKP RKLLVSELAE -1149
45	HSV-Kos	ELDAAAAPGDE PAPPAALPSP AKRPRETPSH ADPPGGASKP RKLLVSELAE -1144
	HSV1-Patton	ELDAAAAPGDE PAPPAALPSP AKRPRETPSP ADPPGGASKP RKLLVSELAE -1144
	HSV1-DJL	ELDAAAAPGDE PAPPAALPSP AKRPRETPSP ADPPGGASKP RKLLVSELAE -1144
	HSV1-F	ELDAAAAPGDE PAPPAALPSP AKRPRETPLH ADPPGGASKP RKLLVSELAE -1144
50	HSV2-MS	DPGYAIARGV PLNTDYYFSH LLGAACVTFK ALFGNNAKIT ESLLKRFIPE -1197
	HSV2-186	DPGYAIARGV PLNTDYYFSH LLGAACVTFK ALFGNNAKIT ESLLKRFIPE -1199
	HSV-Kos	DPAYAIAHGV ALNTDYYFSH LLGAACVTFK ALFGNNAKIT ESLLKRFIPE -1194
	HSV1-Patton	DPAYAIAHGV ALNTDYYFSH LLGAACVTFK ALFGNNAKIT ESLLKRFIPE -1194
	HSV1-DJL	DPAYAIAHGV ALNTDYYFSH LLGAACVTFK ALFGNNAKIT ESLLKRFIPE -1194
55	HSV1-F	DPAYAIAHGV ALNTDYYFSH LLGAACVTFK ALFGNNAKIT ESLLKRFIPE -1194
	HSV2-MS	TWHPPDDVAA RLRAAGFGPA GAGATAEETR RMLHRAFDL A* -1238
	HSV2-186	TWHPPDDVAA RLRAAGFGPA GAGATAEETR RMLHRAFDL A* -1240
	HSV-Kos	VWHPPDDVAA RLRAAGFGAV GAGATAEETR RMLHRAFDL A* -1235
60	HSV1-Patton	VWHPPDDVTA RLRAAGFGAV GAGATAEETR RMLHRAFDL A* -1235
	HSV1-DJL	VWHPPDDVAA RLRTAGFGAV GAGATAEETR RMLHRAFDL A* -1235

Figure 4D Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Aligned by Amino Acid Homology*

5 HSV1-F VWHPPDDVAA RLRAAGFGAV GAGATAEETR RMLHRAFDTL A* -1235

*Amino acid alignment demonstrates difference in amino acid's sequences.

*The gaps “.....” indicate missing amino acids relative to other stanins.

*Wild HSV2-MS is listed as SEQ. ID NO 14.

10 *Wild HSV2-186 is listed as SEQ. ID NO 15.

*Wild HSV-Kos is listed as SEQ. ID NO 16.

*Wild HSV1-Patton is listed as SEQ. ID NO 17.

*Wild HSV1-DJL is listed as SEQ. ID NO 18.

*Wild HSV1-F is listed as SEQ. ID NO 19.

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5 **Figure 5A DNA and amino acid sequence list****SEQ. ID. NO. 1 DNA sequence of DNA polymerase gene for HSV2-MS-M1**

1 ATGTTTGTG CCGCGGGCGG CCCGACTTCC CCCGGGGGGGA AGTCGGCGGC
10 51 TCGGGCGGCG TCTGGGTTTT TTGCCCCCCCA CAACCCCCGG GGAGGCCACCC
101 AGACGGCACC GCCGCCTTGC CGCCGGCAGA ACTTCTACAA CCCCCACCTC
15 151 GCTCAGACCG GAACGCAGCC AAAGGCCCCC GGGCCGGCTC AGCGCCATAC
201 GTACTACAGC GAGTGCAGC AATTTCGATT TATCGCCCCG CGTCGCTGG
251 ACGAGGACGC CCCCGCGGAG CAGCGCACCG GGGTCCACGA CGGCCGCGCTC
301 CGGCGCGCCC CTAAGGTGTA CTGCGGGGGG GACGAGCGCG ACGTCCTCCG
351 CGTGGGCCCCG GAGGGCTTCT GGCCGCGTCG CTTGCGCCTG TGGGGCGGTG
401 CGGACCATGC CCCCAAGGGG TTCGACCCCCA CCGTCACCGT CTTCCACGTG
451 TACGACATCC TGGAGCACGT GGAACACGCG TACAGCATGC GCGCCGCCCCA
501 GCTCCACGAG CGATTTATGG ACGCCATCAC GCCCGCCGGG ACCGTCATCA
551 CGCTTCTGGG TCTGACCCCCC GAAGGCCATC GCGTCGCCGT TCACGTCTAC
601 GGCACGCGGC AGTACTTTA CATGAACAAG GCGGAGGTGG ATCGGCACCT
651 GCAGTGCCGT GCCCCGCGCG ATCTCTGCGA GCGCCTGGCG GCGGCCCTGC
701 GCGAGTCGCC GGGGGCGTCG TTCCGCGCA TCTCCGCGGA CCACTTCGAG
751 GCGGAGGTGG TGGAGCGCGC CGACGTGTAC TATTACGAAA CGCGCCCGAC
801 CCTGTACTAC CGCGTCTTCG TGCGAACGGG GCGCGCGCTG GCCTACCTGT
851 GCGACAACCTT TTGCCCCGCG ATCAGGAAGT ACGAGGGGGG CGTCGACGCC
901 ACCACCCGGT TTATCCTGGA CAACCCGGGG TTTGTCACCT TCGGCTGGTA
951 CCGCCTCAAG CCCGGCGCGG GGAACCGCGCC GGCCCAACCG CGCCCCCCGA
1001 CGGCGTTCGG AACCTCGAGC GACGTCGAGT TTAACTGCAC GCGGACAAC
1051 CTGGCCGTCG AGGGGGCCAT GTGTGACCTG CCGGCCTACA AGCTCATGTG
1101 CTTCGATATC GAATGCAAGG CCGGGGGGGGA GGACGAGCTG GCCTTCCGG
1151 TCGCGGAACG CCCGGAAGAC CTCGTCATCC AGATCTCCTG TCTGCTCTAC
1201 GACCTGTCCA CCACCGCCCT CGAGCACATC CTCCTGTTT CGCTCGGATC

5 **Figure 5B DNA and amino acid sequence list**

1251 CTGCGACCTC CCCGAGTCCC ACCTCAGCGA TCTCGCCTCC AGGGGCCTGC
 1301 CGGCCCGT CGTCCTGGAG TTTGACAGCG AATTGAGAT GCTGCTGGCC
 10 1351 TTCATGACCT TCGTCAAGCA GTACGGCCCC GAGTTCGTGA CCGGGTACAA
 1401 CATCATCAAC TTGACTGGC CCTTCGTCTT GACCAAGCTG ACGGAGATCT
 15 1451 ACAAGGTCCC GCTCGACGGG TACGGCGCA TGAACGGCCG GGGTGTGTC
 1501 CGCGTGTGGG ACATCGGCCA GAGCCACTTT CAGAAGCGCA GCAAGATCAA
 20 1551 GGTGAACGGG ATGGTGAACA TCGACATGTA CGGCATCATC ACCGACAAGG
 1601 TCAAACCTCTC CAGCTACAAG CTGAACGCCG TCGCCGAGGC CGTCTTGAAG
 1651 GACAAGAAGA AGGATCTGAG CTACCGCGAC ATCCCCGCTT ACTACGCCTC
 25 1701 CGGGCCCGCG CAGCGCGGGG TGATCGCGA GTATTGTGTG CAGGACTCGC
 1751 TGCTGGTCGG GCAGCTGTT TCAGTTC TGCCGCACCT GGAGCTTCC
 1801 GCGTCGCGC GCCTGGCGGG CATCAACATC ACCCGCACCA TCTACGACGG
 30 1851 CCAGCAGATC CGCGTCTTCA CGTGCTCTT GCGCCTGCG GGCCAGAAGG
 1901 GCTTCATCCT GCCGGACACC CAGGGCGGT TTGGGGCCT CGACAAGGAG
 35 1951 GCGCCAAGC GCCCGGCCGT GCCTGGGGG GAAGGGGAGC GGCCGGGGGA
 2001 CGGGAACGGG GACGAGGATA AGGACGACGA CGAGGACGAG GACGGGGACG
 2051 AGCGCGAGGA GGTGCGCGC GAGACGGGG GCCGGCACGT TGGGTACCAAG
 40 2101 GGGGCCGGG TCCTCGACCC CACCTCCGGG TTTCACGTCG ACCCGTGGT
 2151 GGTGTTGAC TTTGCCAGCC TGTACCCAG CATCATCCAG GCCCACAAACC
 45 2201 TGTGCTTCAG TACGCTCTCC CTGCGGCCGT AGGCCGTCGC GCACCTGGAG
 2251 GCGGACCGGG ACTACCTGGA GATCGAGGTG GGGGGCCGAC GGCTGTTCTT
 2301 CGTGAAGGCC CACGTACCG AGAGCCTGCT GAGCATCCTG CTGCGCGACT
 50 2351 GGCTGGCCAT GCGAAAGCAG ATCCGCTCGC GGATCCCCA GAGCACCCCC
 2401 GAGGAGGCCG TCCTCCTCGA CAAGCAACAG GCCGCCATCA AGGTGGTGTG
 55 2451 CAACTCGGTG TACGGGTTCA CCAGGGCGCA GCACGGTCTT CTGCCCTGCC
 2501 TGCACCGTGGC CGCCACCGTG ACGACCATCG GCCGCGAGAT GCTCCTCGCG
 2551 ACGCGCGCGT ACGTGCACGC GCGCTGGCG GAGTCGATC AGCTGCTGGC
 60

5 **Figure 5C DNA and amino acid sequence list**

2601 CGACTTCCG GAGGC GGCG GCATGCGCGC CCCC GGTCG TACTCCATGC
10 2651 GCATCATCTA CGGGGACACG GACTCCATT TCGTTTG TG CCGCGGCCTC
2701 ACGGCCGCGG GCCTGGTGGC CATGGCGAC AAGATGGCGA GCCACATCTC
2751 GCGCGCGCTG TTCCCTCCCC CGATCAAGCT CGAGTGGAA AAAACGTTCA
15 2801 CCAAGCTGCT GCTCATCGCC AAGAAAAAGT ACATCGCGT CATCTGCAGG
2851 GGCAAGATGC TCATCAAGGG CGTGGATCTG GTGCGCAAAA ACAACTGCGC
2901 GTTTATCAAC CGCACCTCCA GGGCCCTGGT CGACCTGCTG TTTACGACG
20 2951 ATACCGTATC CGGAGCGGCC GCCGCGTTAG CCGAGCGCCC CGCAGAGGAG
3001 TGGCTGGCGC GACCCCTGCC CGAGGGACTG CAGGC GTTCG GGGCCGTCC
25 3051 CGTAGACGCC CATCGCGCA TCACCGACCC GGAGAGGGAC ATCCAGGACT
3101 TTGTCCCTCAC CGCCGAAC TG AGCAGACACC CGCGCGCGTA CACCAACAAG
3151 CGCCTGGCCC ACCTGACGGT GTATTACAAG CT CATGGCCC GCCGCGCGCA
30 3201 GGTCCCGTCC ATCAAGGACC GGATCCCGTA CGTGATCGTG GCCCAGACCC
3251 GCGAGGTAGA GGAGACGGTC GCGCGGCTGG CCGCCCTCCG CGAGCTAGAC
35 3301 GCGCCGCC CAGGGGACGA GCCCGCCCCC CCAGCGGCC TGCCCTCCCC
3351 GGCCAAGCGC CCCGGGAGA CGCGTCGCA TGCCGACCCC CCGGGAGGCG
3401 CGTCCAAGCC CCGCAAGCTG CTGGTGTCCG AGCTGGCGGA GGATCCCGGG
40 3451 TACGCCATCG CCCGGGCGT TCCGCTAAC ACGGACTATT ACTTCTCGCA
3501 CCTGCTGGGG GCGGCCTGCG TGACGTTCAA GGCCCTGTT GGAAATAACG
45 3551 CCAAGATCAC CGAGAGTCTG TTAAAGAGGT TTATTCCGA GACGTGGCAC
3601 CCCCCGGACG ACGTGGCCGC GCGGCTCAGG GCCGCGGGGT TCGGGCCGGC
3651 GGGGGCCGGC GCTACGGCGG AGGAAACTCG TCGAATGTTG CATAGAGCCT
50 3701 TTGATACTCT AGCATGA

5 **Figure 5D DNA and amino acid sequence list****SEQ. ID. NO. 2 Amino acid sequence of DNA polymerase for HSV2-MS-M1**

1 MFCAAGGPTS PGGKSAARAA SGFFAPHNPR GATQTAPPC RRQNFYNPHL
 10 51 AQTGTQPKAP GPAQRHTYYS ECDEFRFIAP RSLDEDAPAE QRTGVHDGRL
 101 RRAPKVYCGG DERDVLRVGP EGFWPRLRL WGGADHAPKG FDPTVTVFHV
 15 151 YDILEHVEHA YSMRAAQLHE RFMDAITPAG TVITLLGLTP EGHRVAHVY
 201 GTRQYFYMNK AEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE
 251 AEVVERADVY YYETRPTLYY RVFVRSGRAL AYLCDNFCPA IRKYEGGVDA
 301 TTRFILDNPNG FVTFGWYRLK PGRGNAPAQP RPPTAFTGSS DVEFNCTADN
 351 LAVEGAMCDL PAYKLMCFDI ECKAGGEDEL AFPVAERPED LVIQISCLLY
 401 DLSTTALEHI LLPSLGSCDL PESHLSDLAS RGLPAPVVLE FDSEFEMILLA
 451 FMTFKQYGP EFVTGYNIIN FDWPFLVTLKL TEIYKVPLDG YGRMNGRGVF
 501 RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKVKLSSYK LNAVAEAVLK
 551 DKKKDLSYRD IPAYYASGPA QRGVIGEYCV QDSLLVGQLF FKFLPHLELS
 601 AVARLAGINI TRTIYDGQQI RVFTCLLRLA GQKGFLPDT QGRFRGLDKE
 651 APKRPAPVPRG EGERPGDGNG DEDKDDDEDE DGDEREVAR ETGGRHVGYQ
 701 GARVLDPTSG FHVDPVVVFD FASLYPSIIQ AHNLCFSTLS LRPEAVAHLE
 751 ADRDYLEIEV GGRRLLFFVKA HVRESLLSIL LRDWLAMRKQ IRSRIPQSTP
 801 EEAVVLDKQQ AAIKVVCSV YGFTGAQHGL LPCLHVAATV TTIGREMLLA
 851 TRAYVHARWA EFDQLLADFP EAAGMRAPGP YSMRIIYGDT DSIFVLCRGL
 901 TAAGLVAMGD KMASHISRAL FLPPIKLECE KTPTKLLLIA KKKYIGVICG
 951 GKMLIKGVDL VRKNNCAFIN RTSRALVDLL FYDDTVSGAA AALAERPAEE
 1001 WLARPLPEGQ QAFGAFLVDA HRRITDPERD IQDFVLTAE SRHPRAYTNK
 1051 RLAHLTVYYK LMARRAQVPS IKDRIPYVIV AQTRVEETV ARLAALRELD
 1101 AAAPGDEPAP PAALPSPAKR PRETPSHADP PGGASKPRKL LVSELAEDPG
 1151 YAIARGVPLN TDYYFSHLLG AACVTFKALF GNNAKITESL LKRFIPETWH
 1201 PPDDVAARLR AAGFGPAGAG ATAETRRML HRAFDTLA*

5 **Figure 5E DNA and amino acid sequence list****SEQ.ID.NO. 3 DNA sequence of DNA polymerase gene for HSV2-186-M1**

10 1 ATGTTTGTG CCGCGGGCGG CCCGGCTTCC CCCGGGGGGA AGTCGGCGGC
 51 TCGGGCGGCG TCTGGGTTT TTGCCCCCCA CAACCCCCGG GGAGGCCACCC
 101 AGACGGCACC GCCGCCTTGC CGCCGGCAGA ACTTCTACAA CCCCCCACCTC
 151 GCTCAGACCG GAACGCAGCC AAAGGCCCCC GGGCCGGCTC AGCGCCATAC
 201 GTACTACAGC GAGTGCACG AATTGATT TATGCCCG CGTCGCTGG
 251 ACGAGGACGC CCCCGCGGAG CAGCGCACCG GGGTCCACGA CGGCCGCCTC
 301 CGGCGCGCCC CTAAGGTGTA CTGCGGGGGG GACGAGCGCG ACGTCCTCCG
 351 CGTGGGCCCG GAGGGCTTCT GGCGCGTCG CTTGCGCCTG TGGGGCGGTG
 401 CGGACCATGC CCCCAGGGG TTCGACCCCA CCGTCACCGT CTTCCACGTG
 451 TACGACATCC TGGAGCACGT GGAACACGCG TACAGCATGC GCGCCGCCA
 501 GCTCCACGAG CGATTATGG ACGCCATCAC GCCCGCCGGG ACCGTCATCA
 551 CGCTTCTGGG TCTGACCCCA GAAGGCCATC GCGTCGCCGT TCACGTCTAC
 601 GGCACGCGGC AGTACTTTA CATGAACAAG GCGGAGGTGG ATCGGCACCT
 651 GCAGTGCCGT GCCCGCGCG ATCTCTGCGA GCGCCTGGCG GCGGCCCTGC
 701 GCGAGTCGCC GGGGGCGTCG TTCCGCGGA TCTCCGCGGA CCACTTCGAG
 751 GCGGAGGTGG TGGAGCGCGC CGACGTGTAC TATTACGAAA CGCGCCCGAC
 801 CCTGTACTAC CGCGTCTTCG TGCGAAGCGG GCGCGCGCTG GCCTACCTGT
 851 GCGACAACIT TTGCCCGCG ATCAGGAAGT ACGAGGGGGG CGTCGACGCC
 901 ACCACCCGGT TTATCCTGGA CAACCCGGGG TTTGTCACCT TCGGCTGGTA
 951 CCGCCTCAAG CCCGGCCGCG GGAACGCGCC GGCCCAACCG CGCCCCCGA

50 1001 CGGCGTTCGG AACCTCGAGC GACGTCGAGT TTAACTGCAC GGGGGACAAC
 1051 CTGGCCCGTCG AGGGGGCCAT GTGTGACCTG CCGGCCTACA AGCTCATGTG
 1101 CTTCGATATC GAATGCAAGG CCGGGGGGGG GGACGAGCTG GCCTTCCGG
 1151 TCGCGGAACG CCCGGAAGAC CTCGTCATCC AGATCTCCTG TCTGCTCTAC
 1201 GACCTGTCCA CCACCGCCCT CGAGCACATC CTCCGTGTTT CGCTCGGATC

Figure 5F DNA and amino acid sequence list

10 1251 CTGCGACCTC CCCGAGTCCC ACCTCAGCGA TCTCGCCTCC AGGGGCCTGC
 1301 CGGCCCCGT CGTCCTGGAG TTTGACAGCG AATTGAGAT GCTGCTGGCC
 1351 TTCATGACCT TCGTCAAGCA GTACGGCCCC GAGTCGTGA CCGGGTACAA
 15 1401 CATCATCAAC TTGACTGGC CCTTCGTCCT GACCAAGCTG ACGGAGATCT
 1451 ACAAGGTCCC GCTCGACGGG TACGGCGCA TGAACGGCCG GGGTGTGTTC
 20 1501 CGCGTGTGGG ACATCGGCCA GAGCCACTT CAGAACGCGA GCAAGATCAA
 1551 GGTGAACGGG ATGGTGAAACA TCGACATGTA CGGCATCATC ACCGACAAGG
 25 1601 TCAAACCTCTC CAGCTACAAG CTGAACGCCG TCGCCGAGGC CGTCTTGAAG
 1651 GACAAGAAGA AGGATCTGAG CTACCGCGAC ATCCCCGCT ACTACGCC
 1701 CGGGCCCGCG CAGCGCGGGG TGATCGCGA GTATTGTGTG CAGGACTCGC
 30 1751 TGCTGGTCGG GCAGCTGTTT TTCAAGTTTC TGCCGCACCT GGAGCTTCC
 1801 GCCGTCGCGC GCCTGGCGGG CATCAACATC ACCCGCACCA TCTACGACGG
 1851 CCAGCAGATC CGCGTCTTCA CGTGCCTCCT GCGCCTGCG GGCCAGAAGG
 35 1901 GCTTCATCCT GCCGGACACC CAGGGCGGT TTGGGGCGCT CGACAAGGAG
 1951 GCGCCCAAGC GCCCGGCCGT GCCTCGGGGG GAAGGGGAGC GGCCGGGGGA
 40 2001 CGGGAACGGG GACGAGGATA AGGACGACGA CGAGGACGGG GACGAGGACG
 2051 GGGACGAGCG CGAGGAGGTC GCGCGCGAGA CCGGGGGCCG GCACGTTGGG
 45 2101 TACCAGGGGG CCCGGGTCC CGACCCACC TCCGGGTTTC ACGTCGACCC
 2151 CGTGGTGGTG TTGACTTTG CCAGCCTGTA CCCCAGCATC ATCCAGGCC
 2201 AGAACCTGTG CTTGAGTACG CTCTCCCTGC GGGGGAGGC CGTCGCGCAC
 50 2251 CTGGAGGCCGG ACCGGGACTA CCTGGAGATC GAGGTGGGGG GCCGACGGCT
 2301 GTTCTCGTG AAGGCCACG TACCGAGAG CCTGCTGAGC ATCCTGCTGC
 2351 GCGACTGGCT GGCCATGCGA AAGCAGATCC GCTCGCGGAT CCCCCAGAGC
 55 2401 CCCCCCGAGG AGGCCGTCC CCTCGACAAG CAACAGGCCG CCATCAAGGT
 2451 GGTGTGCAAC TCGGTGTACG GGTCACCGG GGCGCAGCAC GGTCTTCTGC
 60 2501 CCTGCCTGCA CGTGGCCGCC ACCGTGACGA CCATCGGCCG CGAGATGCTC

5 **Figure 5G DNA and amino acid sequence list**

2551 CTCGCGACGC GCGCGTACGT GCACGCGCGC TGGGCGGAGT TCGATCAGCT
 10 2601 GCTGGCCGAC TTTCCGGAGG CGGCCGGCAT GCGCGCCCCC GGTCCGTACT
 2651 CCATGCGCAT CATCTACGGG GACACGGACT CCATTTCGT TTTGTGCCGC
 2701 GGCCTCACGG CCGCGGGCCT GGTGGCCATG GGCGACAAGA TGGCGAGCCA
 15 2751 CATCTCGCGC GCGCTGTTCC TCCCCCGAT CAAGCTCGAG TGCGAAAAAA
 2801 CGTTCACCAA GCTGCTGCTC ATCGCCAAGA AAAAGTACAT CGGCGTCATC
 20 2851 TGCGGGGGCA AGATGCTCAT CAAGGGCGTG GATCTGGTGC GCAAAAACAA
 2901 CTGCGCGTTT ATCAACCGCA CCTCCAGGGC CCTGGTCGAC CTGCTGTTT
 2951 ACGACGATAAC CGTATCCGGA GCGGCCGCOG CGTTAGCCGA GCGCCCCGCA
 25 3001 GAGGAGTGGC TGGCGCGACC CCTGCCGAG GGACTGCAGG CGTCGGGGC
 3051 CGTCCTCGTA GACGCCATC GGCGCATCAC CGACCCGGAG AGGGACATCC
 30 3101 AGGACTTTGT CCTCACCGCC GAACTGAGCA GACACCCCGCG CGCGTACACC
 3151 AACAAAGCGCC TGGCCCACCT GACGGTGTAT TACAAGCTCA TGGCCCGCCG
 3201 CGCGCAGGTC CCGTCCATCA AGGACCGGAT CCCGTACGTG ATCGTGGCCC
 35 3251 AGACCCCGGA GGTAGAGGAG ACGGTGCGC GGCTGGCCGC CCTCCGCGAG
 3301 CTAGACGCCG CCGCCCCAGG GGACGAGCCC GCCCCCCCAG CGGCCCTGCC
 40 3351 CTCCCCGGCC AAGCGCCCCC GGGAGACGCC GTCGCATGCC GACCCCCCGG
 3401 GAGGCCGTC CAAGCCCCGC AAGCTGCTGG TGTCCGAGCT GGCGGAGGAT
 3451 CCCGGGTACG CCATGCCCG GGGCGTTCCG CTCAACACGG ACTATTACTT
 45 3501 CTCGCACCTG CTGGGGCGG CCTGCGTGAC GTTCAAGGCC CTGTTGGAA
 3551 ATAACGCCAA GATCACCGAG AGTCTGTTAA AGAGGTTAT TCCCGAGACG
 50 3601 TGGCACCCCC-EGGACGACGT-GGEEEEECEGG-CTCAGGGCGG-EGGGGTTCCG
 3651 GCCGGCGGGG GCCGGCGCTA CGGCGGAGGA AACTCGTCGA ATGTTGCATA
 3701 GAGCCTTGA TACTCTAGCA TGA

5

Figure 5H DNA and amino acid sequence list

SEQ.ID.NO. 4 Amino acid sequence of DNA polymerase for HSV2-186-M1

10	1 MFCAAGGPAS PGGKSAARAA SGFFAPHNPR GATQTAPPPC RRQNFYNPHL 51 AQTGTQPKAP GPAQRHTYYS ECDEFRFIAP RSLDEDAPAE QRTGVHDGRL
15	101 RRAPKVYCGG DERDVLRVGP EGFWPRLRL WGGADHAPEG FDPTVTVFHV 151 YDILEHVEHA YSMRAAQLHE RFMDAITPAG TVITLLGLTP EGHRVAHVY
20	201 GTRQYFYMNK AEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE 251 AEVVERADVY YYETRPTLYY RVFVRSGRAL AYLCDNFPCA IRKYEGGVDA
25	301 TTRFILDNPNG FVTFGWYRLK PGRGNAPAQP RPPTAFGTSS DVEFNCTADN 351 LAVEGAMCDL PAYKLMCFDI ECKAGGEDEL AFPVAERPED LVIQISCLLY
30	401 DLSTTALEHI LLFSLGSCDL PESHLSDLAS RGLPAPVVLE FDSEFEMILLA 451 FMTFKVKQYGP EFVTGYNIIN FDWPFLTKL TEIYKVPLDG YGRMNGRGVF
35	501 RVWDIGQSHF QKRSKIKVNG MVNIDMYGII TDKVKLSSYK LNAVAEAVLK 551 DKKKDLSYRD IPAYYYAS GPA QRGVIGEYCV QDSLLVGQLF FKFLPHLELS
40	601 AVARLAGINI TRTIYDGQQI RVFTCLLRLA GQKGFLPDT QGRFRGLDKE 651 APKRPAVPRG EGERPGDGNG DEDKDDDEDG DEDGDREEV ARETGGRHVG
45	701 YQGARVLDPT SGFHVDPVVV FDFASLYPSI IQAHNLCFST LSLRPEAVAH 751 LEADRDXLEI EVGGRRLLFFV KAHVRESLLS ILLRDWLAMR KQIRSRIQPS
50	801 PPEEAVIDDK QQAAIKVVCN SVYGFTGAQH GLLPCLHVAA TVTTIGREML 851 LATRAYVHAR WAEFDQLLAD FPEAAGMRAP GPYSMIRIYG DTDSIFVLCR
55	901 GLTAAGLVAM GDKMASHISR ALFLPPIKLE CEKTPTKLLL IAKKKYIGVI 951 CGGKMLIKGV DLVRKNNCAC INRTSRALVD LLFYDDTVSG AAAALAERPA
60	1001 EEWLARPLPE GLQAFGAVLV DAHRRITDPE RDIQDFVLT A ELSRHPRAYT 1051 NKRLAHLT VY YKLMARRAQV PSIKDRIPYV IVAQTREVEE TVARLAALRE 1101 LDAAA PGD EPP APPAALPSA KRPRETPSHA DPPGGASKPR KLLVSELAED 1151 PGYAIARGVP LNTDYYFSHL LGAACVTFKA LFGNNAKITE SLLKRFIPET 1201 WHPPDDVAAR LRAAGFGPAG AGATAEEERR MLHRAFDTLA *

5 **Figure 5I DNA and amino acid sequence list****SEQ.ID.NO. 5 DNA sequence of DNA polymerase gene for HSV1-KOS-M1**

10 1 ATGTTTCCG GTGGCGCGG CCCGCTGTCC CCCGGAGGAA AGTCGGCGC
 51 CAGGGCGCG TCCGGGTTT TTGCGCCCGC CGGCCCTCGC GGAGCCGGCC
 101 GGGGACCCCC GCCTTGTTG AGGCAAAACT TTTACAACCC CTACCTCGCC
 15 151 CCAGTCGGGA CGCAACAGAA GCCGACCGGG CCAACCCAGC GCCATACGTA
 201 CTATAGCGAA TGCGATGAAT TTCGATTAT CGCCCCGCGG GTGCTGGACG
 20 251 AGGATGCCCG CCCGGAGAAG CGCGCGGGG TGCACGACGG TCACCTCAAG
 301 CGCGCCCCA AGGTGTACTG CGGGGGGGAC GAGCGCGACG TCCTCCGCGT
 351 CGGGTCGGGC GGCTTCTGGC CGCGCGCTC GCGCCTGTGG GGCGCGTGG
 25 401 ACCACGCCCG GGCGGGGTTA AACCCCACCG TCACCGTCTT TCACGTGTAC
 451 GACATCCTGG AGAACGTGGA GCACGCGTAC GGCATGCGCG CGGCCAGTT
 30 501 CCACGCGCGG TTTATGGACG CCATCACACC GACGGGGACC GTCATCACGC
 551 TCCTGGGCCT GACTCCGGAA GGCCACCGGG TGGCCGTTCA CGTTTACGGC
 601 ACGCGGCAGT ACTTTACAT GAACAAGGAG GAGGTTGACA GGCACCTACA
 35 651 ATGCCGCGCC CCACGAGATC TCTGCGAGCG CATGGCCGCG GCCCTGCGCG
 701 AGTCCCCGGG CGCGTCGTTT CGCGGCATCT CCGCGGACCA CTTCGAGGCG
 40 751 GAGGTGGTGG AGCGCACCGA CGTGTACTAC TACGAGACGC GCCCCGCTCT
 801 GTTTACCGC GTCTACGTCC GAAGCGGGCG CGTGCTGTGCG TACCTGTGCG
 851 ACAACTCTG CCCGGCCATC AAGAAGTACG AGGGTGGGGT CGACGCCACC
 45 901 ACCCGGTTCA TCCTGGACAA CCCCGGGTTT GTCACCTTCG GCTGGTACCG
 951 TCTCAAACCG GGCGGAAACA ACACGCTAGC CCAGCCGCGG GCCCCGATGG
 50 1001 CCTTCGGGAC ATCCAGCGAC GTCGAGTTA ACTGTACGGC GGACAACCTG
 1051 GCCATCGAGG GGGGCATGAG CGACCTACCG GCATACAAGC TCATGTGCTT
 1101 CGATATCGAA TGCAAGGCAGG GGGGGGAGGA CGAGCTGGCC TTTCCGGTGG
 55 1151 CGGGGCACCC GGAGGACCTG GTTATTAGA TATCCTGTCT GCTCTACGAC
 1201 CTGTCCACCA CCGCCCTGGA GCACGTCCTC CTGTTTCGC TCGGTTCTG

5

Figure 5J DNA and amino acid sequence list

10 1251 CGACCTCCCC GAATCCCACC TGAACGAGCT GGCGGCCAGG GGCTGCCCA
 1301 CGCCCGTGGT TCTGGAATT CACAGCGAAT TCGAGATGCT GTTGGCCTTC
 1351 ATGACCCTTG TGAAACAGTA CGGCCCGAG TTCGTGACCG GGTACAACAT
 15 1401 CATCAACTTC GACTGGCCCT TCTTGCTGGC CAAGTTGACG GACATTACA
 1451 AGGTCCCCCT GGACGGGTAC GGCGCATGA ACGGCCGGGG CGTGTTCGC
 20 1501 GTGTGGGACA TAGGCCAGAG CCACTTCCAG AAGCGCAGCA AGATAAAGGT
 1551 GAACGGCATG GTAACATCG ACATGTACGG GATCATAACC GACAAGATCA
 1601 AGCTCTCGAG CTACAAGCTC AACGCCGTGG CCGAAGCCGT CCTGAAGGAC
 25 1651 AAGAAGAAGG ACCTGAGCTA TCGCGACATC CCCGCCTACT ACGCCGCCGG
 1701 GCCCGCGCAA CGCGGGGTGA TCGCGAGTA CTGCATACAG GATTCCCTGC
 30 1751 TGGTGGGCCA GCTTTTTT AAGTTTTGC CCCATCTGGA GCTCTCGGCC
 1801 GTCGCGCGCT TGGCGGGTAT TAACATCACC CGCACCATCT ACGACGGCCA
 1851 GCAGATCCGC GTCTTACGT GCCTGCTGCG CCTGGCCGAC CAGAAGGGCT
 35 1901 TTATTCTGCC GGACACCCAG GGGCGATTAA GGGGCGCCGG GGGGGAGGCG
 1951 CCCAAGCGTC CGGCCGCAGC CCGGGAGGAC GAGGAGCGC CAGAGGAGGA
 40 2001 GGGGGAGGAC GAGGACGAAC GCGAGGAGGG CGGGGGCGAG CGGGAGCCGG
 2051 AGGGCGCGCG GGAGACCGCC GGCCGGCACG TGGGGTACCA GGGGGCCAGG
 2101 GTCCTTGACC CCACCTCCGG GTTTCACGTG AACCCCGTGG TGGTGTTCGA
 45 2151 CTTGCCAGC CTGTACCCCA GCATCATCCA GGCCCACAAAC CTGTGCTTCA
 2201 GCACGGCTCTC CCTGAGGGCC GACCGAGTGG CGCACCTGGA GGGGGGCAAG
 2251 GACTACCTGG AGATCGAGGT GGGGGGGCGA CGGCTTTCT TCGTCAAGGC
 2301 TCACGTGCGA GAGAGCCTCC TCAGCATCCT CCTGCGGGAC TGGCTCGCCA
 2351 TGCAGAAAGCA GATCCGCTCG CGGATTCCCC AGAGCAGCCC CGAGGAGGCC
 55 2401 GTGCTCCTGG ACAAGCAGCA GGCCGCCATC AAGGTCGTGT GTAACTCGGT
 2451 GTACGGGTTA ACAGGGAGCGC AGCACGGACT CCTGCCGTGC CTGCACGTTG
 60 2501 CCGCGACGGT GACGACCATC GGCCGCGAGA TGCTGCTCGC GACCCGCGAG

5 **Figure 5K DNA and amino acid sequence list**

2551 TACGTCCACG CGCGCTGGGC GGCCTTCGAA CAGCTCCTGG CCGATTTCCC
10 2601 GGAGGCAGGCC GACATGCGCG CCCCCGGGCC CTATTCCATG CGCATCATCT
2651 ACGGGGACAC GGACTCCATA TTTGTGCTGT GCCGCGGCCT CACGGCCGCC
15 2701 GGGCTGACGG CCATGGGCGA CAAGATGGCG AGCCACATCT CGCGCGCGCT
2751 GTTTCTGCC CCCATCAAAC TCGAGTGCAG AAAGACGTT ACCAAGCTGC
2801 TGCTGATCGC CAAGAAAAAG TACATCGCG TCATCTACGG GGGTAAGATG
20 2851 CTCATCAAGG GCGTGGATCT GGTGCGCAAA ACAACTGCG CGTTTATCAA
2901 CGCGCACCTCC AGGGCCCTGG TCGACCTGCT GTTTACGAC GATACCGTAT
2951 CCGGAGCGGC CGCCGCGTTA GCGAGCGCC CCGCAGAGGA GTGGCTGGCG
25 3001 CGACCCCTGC CCGAGGGACT GCAGGCGTTC GGGGCCGTCC TCGTAGACGC
3051 CCATCGGCGC ATCACCGACC CGGAGAGGGA CATCCAGGAC TTTGTCCCTCA
30 3101 CCGCCGAACG GAGCAGACAC CCGCGCGCGT ACACCAACAA GCGCCTGGCC
3151 CACCTGACGG TGTATTACAA GCTCATGGCC CGCCGCGCGC AGGTCCCGTC
3201 CATCAAGGAC CGGATCCCGT ACGTGATCGT GGCCCAGACC CGCGAGGTAG
35 3251 AGGAGACGGT CGCGCGGCTG GCCGCCCTCC GCGAGCTAGA CGCCGCCGCC
3301 CCAGGGGACG AGCCCGCCCC CCCCAGGCC CTGCCCTCCC CGGCCAAGCG
40 3351 CCCCCGGGAG ACGCCGTGCG ATGCCGACCC CCCGGGAGGC GCGTCCAAGC
3401 CCCGCAAGCT GCTGGTGTCC GAGCTGGCCG AGGATCCCGC ATACGCCATT
3451 GCCCACGGCG TCGCCCTGAA CACGGACTAT TACTTCTCCC ACCTGTTGGG
45 3501 GGCGGCGTGC GTGACATTCA AGGCCCTGTT TGGGAATAAC GCCAAGATCA
3551 CCGAGAGTCT GTAAAAAAGG TTTATTCCCG AAGTGTGGCA CCCCCCGGAC
50 3601 GACGTGGCCG CGCGGCTCCG GGCGCGAGGG TTCGGGGCGG TGGGTGCCGG
3651 CGCTACGGCG GAGGAAACTC GTGAAATGTT GCATAGAGCC TTTGATACTC
55 3701 TAGCATGA

5 **Figure 5L DNA and amino acid sequence list**

SEQ.ID.NO. 6 Amino acid sequence of DNA polymerase for HSV1-KOS-M1

1 MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGRGPPLC RQNFYNPYLA
 10 51 PVGTQQKPTG PTQRHTYYSE CDEFRFIAPR VLDEDAPPEK RAGVHDGHLK
 101 RAPKVYCGGD ERDVLRVGSG GFWPRRSRLW GGVDHAPAGF NPTVTVFHVY
 15 151 DILENVEHAY GMRAAQPHAR FMDAITPTGT VITLLGLTPE GHRVAVHVG
 201 201 TRQYFYMNKE EVDRHLQCRA PRDLCEMAA ALRESPGASF RGISADHFEA
 251 251 EVVERTDVYY YETRPALFYR VYVRSGRVL S YLCDNFCPAI KKYEGGVDAT
 301 301 TRFILDNPFG VTFGWYRLKP GRNNTLAQPR APMAFGTSSD VEFNCTADNL
 351 351 AIEGGMSDLP AYKLMCFDIE CKAGGEDELA FPVAGHPEDL VIQISCLLYD
 401 401 LSTTALEHVL LP SLGSCDLP ESHLNELAAR GLPTPVVLEF DSEFEMLLAF
 451 451 MTLVKQYGPE FVTGYNIINF DWPFLAKLT DIYKVPLDGY GRMNGRGVFR
 501 501 VWDIGQSHFQ KR SKIKVNGM VNIDMYGIIT DKIKLSSYKL NAVA EAVLKD
 551 551 KKKDLSYRDI PAYYAAGPAQ RGVIGEYCIQ DSLLVGQLFF KFLPHLELSA
 601 601 VARLAGINIT RTTYDGQQIR VFTCLLRLAD QKGFI LPDTQ GRFRGAGGEA
 651 651 PKRPA AARED EERPEEEGED EDEREEGGGE REPEGARETA GRHVGYQGAR
 701 701 VLDPTSGFH V NPVVVFDFAS LYP SIIQAHN LCFSTLSLR A DAVA HLEAGK
 751 751 DYLEIEVGGR RLFFVKAHVR ESLLSILLRD WLAMRKQIRS RIPQSSPEEA
 801 801 VLLDKQQAAI KVVCNSVYGF TGAQHGLLPC LHVAATVTTI GREMLLATRE
 851 851 YVHARWAAFE QLLADFP EAA DMRAPGPYSM RIYGDTSI FVLCRGLTAA
 901 901 GLTAMGDKMA SHISRALFLP PIKLECEKTF TKLLLIAKKK YIGVITYGGKM
 951 951 LIKGVDLVRK NNCAFINRTS RALVDLLFYD DTVSGAAAAL AERPAEEWLA
 1001 1001 RPLPEGLQAF GAVLVDAHRR ITDPERDIQD FVLTAELSRH PRAYTNKRLA
 1051 50 1051 HLT VYYKLMA RRAQVPSIKD RIPYVIVAQT REVEETVARL AALRELDAAA
 1101 1101 PGDEPAPPAA LPSPA KRPRE TPSHADPPGG ASKPRKLLVS ELAEDPAYAI
 1151 55 1151 AHGVALNTDY YFSHLLGAAC VTFKALFGNN AKITESLLKR FIPEVWHPPD
 1201 1201 DVAARLRAAG FGAVGAGATA EETRRMLHRA FDTLA*

5 **Figure 5M DNA and amino acid sequence list****SEQ.ID.NO. 7 DNA sequence of HSV polymerase gene for HSV1-F-M1**

10	1	ATGTTTCCG GTGGCGCGG CCCGCTGTCC CCCGGAGGAA AGTCGGCGC
	51	CAGGGCGCG TCCGGGTTTT TTGCGCCCGC CGGCCCTCGC GGAGCCGGCC
	101	GGGGACCCCC GCCTTGCTTG AGGCAAAACT TTTACAACCC CTACCTCGCC
15	151	CCAGTCGGGA CGCAACAGAA GCCGACCGGG CCAACCCAGC GCCATACGTA
	201	CTATAGCGAA TGCGATGAAT TTGCGATTCA CGCCCCCGGG GTGCTGGACG
20	251	AGGATGCCCG CCCGGAGAAG CGCGCCGGG TGCACGACGG TCACCTCAAG
	301	CGCGCCCCCA AGGTGTACTG CGGGGGGGAC GAGCGCGACG TCCTCCGCGT
	351	CGGGTCGGGC GGCTTCTGGC CGCGCGCCTC GCGCCTGTGG GCGGCGTGG
25	401	ACCACGCCCGG GGCGGGGTTTC AACCCCACCG TCACCGTCTT TCACGTGTAC
	451	GACATCCTGG AGAACGTGGA GCACGCGTAC GGCATGCGCG CGGCCAGTT
30	501	CCACGCGCGG TTTATGGACG CCATCACACC GACGGGGACC GTCATCACGC
	551	TCCTGGGCCT GACTCCGGAA GGCCACCGGG TGGCCGTTCA CGTTTACGGC
	601	ACGCGGCAGT ACTTTACAT GAACAAGGAG GAGGTGACGAC GGCACCTACA
35	651	ATGCCCGCGCC CCACGAGATC TCTGCGAGCG CATGGCCGCG GCCCTGCGCG
	701	AGTCCCCGGG CGCGTCGTTTC CGCGGCATTT CCGCGGACCA CTTCGAGGCG
40	751	GAGGTGGTGG AGCGCACCGA CGTGTACTAC TACGAGACGC GCCCCGCTCT
	801	GTTTTACCGC GTCTACGTCC GAAGCGGGCG CGTGCTGTGCG TACCTGTGCG
	851	ACAACCTCTG CCCGGCCATC AAGAAGTACG AGGGTGGGGT CGACGCCACC
45	901	ACCCGGTTCA TCCTGGACAA CCCGGGGTTTC GTCACCTTCG GCTGGTACCG
	951	TCTCAAACCG GGCGGAACA ACACGCTAGC CCAGCCGCGG GCCCCGATGG
	1001	CCTTCGGGAC ATCCAGCGAC GTCGAGTTA ACTGTACGGC GGACAACCTG
50	1051	GCCATCGAGG GGGGCATGAG CGACCTACCG GCATACAAGC TCATGTGCTT
	1101	CGATATCGAA TGCAAGGCGG GGGGGGAGGA CGAGCTGGCC TTTCCGGTGG
55	1151	CCGGGCACCC GGAGGACCTG GTCATCCAGA TATCCTGTCT GCTCTACGAC
	1201	CTGTCCACCA CCGCCCTGGA GCACGTCCTC CTGTTTCGC TCGGTTCCCTG
	1251	CGACCTCCCC GAATCCCACC TGAACGAGCT GGCGGCCAGG GGCCTGCCA
60		

5 **Figure 5N DNA and amino acid sequence list**

	1301	CGCCCGTGGT TCTGGAATTC GACAGCGAAT TCGAGATGCT GTTGGCCTTC
10	1351	ATGACCCCTTG TGAAACAGTA CGGCCCGAG TTCTGTGACCG GGTACAACAT
	1401	CATCAACTTC GACTGGCCCT TCTTGCTGGC CAAGCTGACG GACATTACAA
	1451	AGGTCCCCCT GGACGGGTAC GGCGCGATGA ACGGCCCCGG CGTGTTCGC
15	1501	GTGTGGGACA TAGGCCAGAG CCACCTCCAG AAGCGCAGCA AGATAAAGGT
	1551	GAACGGCATG GTGAACATCG ACATGTACGG GATTATAACC GACAAGATCA
20	1601	AGCTCTCGAG CTACAAGCTC AACGCCGTGG CCGAAGCCGT CCTGAAGGAC
	1651	AAGAAGAAGG ACCTGAGCTA TCGCGACATC CCCGCCTACT ACGCCGCCGG
	1701	GCCCGCGCAA CGCGGGGTGA TCGGCGAGTA CTGCATACAG GATTCCCTGC
25	1751	TGGTGGGCCA GCTGTTTTTAAAGTTTTGC CCCATCTGGA GCTCTCGGCC
	1801	GTCGCGCGCT TGGCGGGTAT TAACATCACC CGCACCATCT ACGACGGCCA
30	1851	GCAGATCCGC GTCTTTACGT GCCTGCTGCG CCTGGCCGAC CAGAAGGGCT
	1901	TTATTCTGCC GGACACCCAG GGGCGATTAA GGGGCGGCCGG GGGGGAGGCG
	1951	CCCAAGCGTC CGGCCGCAGC CCGGGAGGAC GAGGAGCGGC CAGAGGAGGA
35	2001	GGGGGAGGAC GAGGACGAAC GCGAGGAGGG CGGGGGCGAG CGGGAGCCGG
	2051	AGGGCGCGCG GGAGACCGCC GGCGGCACG TGGGGTACCA GGGGGCCAGG
40	2101	GTCCTTGACC CCACCTCCGG GTTTCATGTG AACCCCGTGG TGGTGTTCGA
	2151	CTTTGCCAGC CTGTACCCCA GCATCATCCA GGCCCACAAC CTGTGCTTC
	2201	GCACGCTCTC CCTGAGGGCC GACGCAGTGG CGCACCTGGA GGCAGGGCAAG
45	2251	GACTACCTGG AGATCGAGGT GGGGGGGCGA CGGCTGTTCT TCGTCAAGGC
	2301	TCACGTGCGA GAGAGCCTCC TCAGCATCCT CCTGCGGGAC TGGCTCGCCA
	2351	TGCGAAAGCA GATCCGCTCG CGGATTCCCC AGAGCAGCCC CGAGGAGGCC
50	2401	GTGCTCCTGG ACAAGCAGCA GGCGCCATC AAGGTCGTGT GTAACCTCGGT
	2451	TTACGGGTTAC CGGGGAGCGC AGCACGGACT CCTGCCGTGC CTGCACGTTG
55	2501	CCGCGACGGT GACGACCATC GGCGCGAGA TGCTGCTCGC GACCCGCGAG
	2551	TACGTCCACG CGCGCTGGGC GGCGTTGAA CAGCTCCTGG CCGATTTC
60	2601	GGAGGCGGCC GACATGCGCG CCCCCGGGCC CTATTCCATG CGCATCATCT

Figure 50 DNA and amino acid sequence list

10	2651	ACGGGGACAC GGACTCCATC TTTGTGCTGT GCCGCGGCCT CACGGCCGCC
	2701	GGGCTGACGG CCGTGGCGA CAAGATGGCG AGCCACATCT CGCGCGCGCT
	2751	GTTTCTGTCC CCCATCAAAC TCGAGTGCAG AAAGACGTT ACCAAGCTGC
15	2801	TGCTGATCGC CAAGAAAAAG TACATCGGCG TCATCTACGG GGGTAAGATG
	2851	CTCATCAAGG GCGTGGATCT GGTGCGCAAA AACAACTGCG CGTTTATCAA
20	2901	CCGCACCTCC AGGGCCCTGG TCGACCTGCT GTTTTACGAC GATACCGTAT
	2951	CCGGAGCGGC CGCCGCGTTA GCCGAGCGCC CCGCAGAGGA GTGGCTGGCG
	3001	CGACCCCTGC CCGAGGGACT GCAGGCGTTC GGGGCCGTCC TCGTAGACGC
25	3051	CCATCGGCGC ATCACCGACC CGGAGAGGGGA CATCCAGGAC TTTGTCCCTCA
	3101	CCGCCGAACT GAGCAGACAC CCGCGCGCGT ACACCAAACAA GCGCCTGGCC
30	3151	CACCTGACGG TGTATTACAA GCTCATGGCC CGCCGCGCGC AGGTCCCCGTC
	3201	CATCAAGGAC CGGATCCCCT ACGTGATCGT GGCCCAGACC CGCGAGGTTAG
	3251	AGGAGACGGT CGCGCGGCTG GCCGCCCTCC GCGAGCTCGA CGCCGCGGCC
35	3301	CCAGGGGACG AGCCCCCCCC CCCCCGCGGCC CTGCCCTCCCC CGGCCAAGCG
	3351	CCCCCGGGAG ACGCCGTTGC ATGCCGACCC CCCGGGAGGC GCGTCCAAGC
40	3401	CCCGCAAGCT GCTGGTGTCC GAGCTGGCCG AGGATCCCGC ATACGCCATT
	3451	GCCCCACGGCG TCGCCCTGAA CACGGACTAT TACTTCTCCC ACCTGTTGGG
	3501	GGCGCGGTGC GTGACATTCA AGGCCCTGTT TGGGAATAAC GCCAAGATCA
45	3551	CCGAGAGTCT GTTAAAAAGG TTTATTCCCG AAGTGTGGCA CCCCCCGGAC
	3601	GACGTGGCCG CGCGGCTCCG GCCCGCAGGG TTCGGGGCGG TGGGTGCCGG
50	3651	CGCTACGGCG GAGGAAACTC GTCGAATGTT GCATAGAGCC TTTGATACTC
	3701	TAGCATGA

5 **Figure 5P DNA and amino acid sequence list**

SEQ.ID.NO. 8 Amino acid sequence of DNA polymerase for HSV1-F-M1

10	1 MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGRGPPLC RQNFYNPYLA 51 PVGTQQKPTG PTQRHTYYSE CDEFRFLAPR VLDEDAPPEK RAGVHDGHLK
15	101 RAPKVYCGGD ERDVLRVGSG GFWPRRSRLW GGVDHAPAGF NPTVTVFHVY 151 DILENVEHAY GMRAAQFHAR FMDAITPTGT VITLLGLTPE GHRVAVHVYG
20	201 TRQYFYMNKE EVDRHLQCRA PRDLCEMAA ALRESPGASF RGISADHFEA 251 EVVERTDVYY YETRPALFYR VYVRSGRVLS YLCDNFCPAI KKYEGGVDAT
25	301 TRFILDNPFGF VTFGWYRLKP GRNNTLAQPR APMAFGTSSD VEFNCTADNL 351 AIEGGMSDLP AYKLMCFDIE CKAGGEDELA FPVAGHPEDL VIQISCLLYD
30	401 LSTTALEHVL LFSLGSCDLP ESHLNELAAR GLPTPVVLEF DSEFEMLLAF 451 MTLVKQYGPE FVTGYNIINF DWPFLLAKL DTIYKVPLDGY GRMNGRGVFR
35	501 VWDIGQSHFQ KRSKIKVNGM VNIDMYGIIT DKIKLSSYKL NAVAEAVLKD 551 KKKDLSYRDI PAYYAAGPAQ RGVIGEYCIQ DSLLVGQLFP KFLPHILESA
40	601 VARLAGINIT RTIYDGQQIR VFTCLLRLAD QKGFILPDTQ GRFRGGGEA 651 PKRPAAARED EERPEEEGED EDEREEGGGE REPEGARETA GRHVGYQGAR
45	701 VLDPTSGFHV NPVVVFDFAS LYPHIQAHN LCFSTLSLRA DAVAHLEAGK 751 DYLIIEVGGR RLFFVKAHVR ESLLSILLRD WLAMRKQIRS RIPQSSPEEA
50	801 VLLDKQQAAI KVVCNSVYGF TGAQHGLLPC LHVAATVTTI GREMLLATRE 851 YVHARWAAFE QLLADPPEAA DMRAPGPYSM RIYGDTSI FVLCRGLTAA
55	901 GLTAVGDKMA SHISRALFLS PIKLECEKTF TKLLLIAKKK YIGVITYGGKM 951 LIKGVDLVRK NNCAFINRTS RALV DLLFYD DTVSGAAAAL AERPAEEWLA
60	1001 RPLPEGLQAF GAVLVDAHRR ITDPERDIQD FVLTAELSRH PRAYTNKRLA 1051 HLTIVYYKLMA RRAQVPSIKD RIPYVIVQAQT REVEETVARL AALRELDAAA 1101 PGDEPAPPAA LPSPAKRPRE TPLHADPPGG ASKPRKLLVS ELAEDPAYAI 1151 AHGVALNTDY YFSHLLGAAC VTFKALFGNN AKITESLLKR FIPEVWHPPD 1201 DVAARLRAAG FGAVGAGATA EETRRMLHRA FDTLA*

5 **Figure 5Q DNA and amino acid sequence list****SEQ.ID.NO. 9 DNA sequence of HSV polymerase gene for HSV1-DJL-M1**

1 ATGTTTCGG GTGGCGCGG CCCGCTGTCC CCCGGAGGAA AGTCGGCGGC
 10 51 CAGGGCGCG TCCGGGTTT TTGCGCCCGC CGGCCCTCGC GGAGCCGGCC
 101 GGGGACCCCC GCCTTGTGG AGGCAAAACT TTTACAACCC CTACCTCGCC
 15 151 CCAGTCGGGA CGCAACAGAA GCCGACCGGG CCAACCCAGC GCCATACGTA
 20 201 CTATAGCGAA TGCGATGAAT TTCGATTAT CGCCCCGCGG GTGCTGGACG
 25 251 AGGATGCCCG CCCGGAGAAG CGCGCCGGGG TGACGACGG TCACCTCAAG
 30 301 CGCGCCCCCA AGGTGTACTG CGGGGGGGAC GAGCGCGACG TCCTCCGCGT
 35 351 CGGGTCGGGC GGCTTCTGGC CGCGCGCTC GCGCCTGTGG GGCGGCGTGG
 40 401 ACCACGCCCGG GGCGGGGTTTC AACCCCACCG TCACCGTCTT TCACGTGTAT
 45 451 GACATCCTGG AGAACGTGGA GCACCGTAC GGCAATGCGCG CGGCCAGTT
 50 501 CCACGCGCGG TTTATGGACG CCATCACACC GACGGGGACC GTCATCACGC
 55 551 TCCTGGGCCT GACTCCGGAA GGCCACCGGG TGGCCGTTCA CGTTTACGGC
 60 601 ACGCGGCAGT ACTTTACAT GAACAAGGAG GAGGTTGACA GGCACCTACA
 65 651 ATGCCGCGCC CCACGAGATC TCTGCGAGCG CATGGCCGCG GCCCTGCGCG
 70 701 AGTCCCCGGG CGCGTCGTTTC CGCGGCATCT CCGCGGACCA CTTCGAGGCG
 75 751 GAGGTGGTGG AGCGCACCGA CGTGTACTAC TACGAGACGC GCCCCGCTCT
 80 801 GTTTTACCGC GTCTACGTCC GAAGCGGGCG CGTGCTGTGCG TACCTGTGCG
 85 851 ACAACTTCTG CCCGGCCATC AAGAAGTACG AGGGTGGGGT CGACGCCACC
 90 901 ACCCGGTCA TCCTGGACAA CCCCGGGTTTC GTCACCTTCG GCTGGTACCG
 95 951 TCTCAAACCG GGCGGAACA ACACGCTAGC CCAGCCGCGG GCCCCGATGG
 100 1001 CCTTCGGGAC ATCCAGCGAT GTCGAGTTA ACTGTACGGC GGACAACCTG
 105 1051 GCCATCGAGG GGGGCATGAG CGACCTACCG GCATACAAGC TCATGTGCTT
 110 1101 CGATATCGAA TGCAAGGCGG GGGGGGAGGA CGAGCTGGCC TTTCCGGTGG
 115 1151 CGGGGCACCC GGAGGACCTG GTCATCCAGA TATCCTGTCT GCTCTACGAC
 120 1201 CTGTCCACCA CCGCCCTGGA GCACGTCTC CTGTTTCGC TCGGTTCTG
 125 1251 CGACCTCCCC GAATCCCACC TGAACGAGCT GGCGGCCAGG GGCTGCCCA

Figure 5R DNA and amino acid sequence list

10 1301 CGCCCGTGGT TCTGGAATT GACAGCGAAT TCGAGATGCT GTTGGCCTTC
1351 ATGACCCTTG TGAAACAGTA CGGCCCGAG TTCTGTACCG GGTACAACAT
1401 AATCAACTTC GACTGGCCCT TCTTGCTGGC CAAGCTGACG GACATTTACA
1451 AGGTCCCCCT GGACGGGTAC GGCGCATGA ACGGCCGGGG CGTGTTCGC
1501 GTGTGGGACA TAGGCCAGAG CCACTTCCAG AAGCGCAGCA AGATAAAGGT
1551 GAACGGCATG GTGAACATCG ACATGTACGG GATTATAACC GACAAGATCA
20 1601 AGCTCTCGAG CTACAAGCTC AACGCCGTGG CCGAAGCCGT CCTGAAGGAC
1651 AAGAAGAAGG ACCTGAGCTA TCGCGACATC CCCACCTACT ACGCCGCCGG
25 1701 GCCCGCGCAA CGCGGGGTGA TCGCGAGTA CTGCATACAG GATTCCCTGC
1751 TGGTGGGCCA GCTGTTTTT AAGTTTTGC CCCATCTGGA GCTCTGGCC
30 1801 GTCGCGCGCT TGGCGGGTAT TAACATCACC CGCACCATCT ACGACGGCCA
1851 GCAGATCCGC GTCTTACGT GCCTGCTGCG CCTGGCCGAC CAGAAGGGCT
1901 TTATTCTGCC GGACACCCAG GGGCGATTAA GGGGCGCCGG GGGGGAGGCG
35 1951 CCCAAGCGTC CGGCCGCAGC CGGGGAGGAC GAGGAGCGGC CAGAGGAGGA
2001 GGGGGAGGAC GAGAACGAAC GCGAGGAGGG CGGGGGCGAG CGGGAGCCGG
40 2051 AGGGCGCGCG GGAGACCGCC GGCCGGCAAG TGGGGTACCA GGGGGCCAGG
2101 GTCCTTGACC CCACCTCCGG GTTTCACGTG AACCCCGTGG TGGTGTTCGA
2151 CTTTGCCAGC CTGTACCCCA GCATCATCCA GGCCCACAAAC CTGTGCTTCA
45 2201 GCACGCTCTC CCTGAGGGCC GACGCAGTGG CGCACCTGGA GGCGGGCAAG
2251 GACTACCTGG AGATCGAGGT GGGGGGGCGA CGGCTGTTCT TCGTCAAGGC
50 2301 TCACGTGCGA GAGAGCCTCC TCAGCATCCT CCTGCGGGAC TGGCTCGCCA
2351 TCGGAAAGCA GATCCGCTCG CGGATTCCCC AGAGCAGCCC CGAGGAGGCC
2401 GTGCTCCTGG ACAAGCAGCA GGCCGCCATC AAGGTCGTGT GTAACTCGGT
55 2451 TTACGGGTTACCGGGAGCGC AGCACGGACT CCTGCCGTGC CTGCACGTTG
2501 CCGCGACGGT GACGACCATC GGCGCGAGA TGCTGCTCGC GACCCGCGAG
2551 TACGTCCACG CGCGCTGGGC GGCCTTCGAA CAGCTCCTGG CCGATTCCC

5 **Figure 5S DNA and amino acid sequence list**

2601 GGAGGCGGCC GACATGCGCG CCCCCGGGCC CTATTCCATG CGCATCATCT
10 2651 ACGGGGACAC GGACTCCATA TTTGTGCTGT GCCGCGGCCT CACGGCCGCC
 2701 GGGCTGACGG CCGTGGGCGA CAAGATGGCG AGCCACATCT CGCGCGCGCT
15 2751 GTTCTGCC CCCATCAAAC TCGAGTGCAG AAAGACGTTCA ACCAAGCTGC
 2801 TGCTGATCGC CAAGAAAAAG TACATCGGCG TCATCTACGG GGGTAAGATG
 2851 CTCATCAAGG GCGTGGATCT GGTGCGCAAA AACAACTGCG CGTTTATCAA
20 2901 CCGCACCTCC AGGGCCCTGG TCGACCTGCT GTTTACGAC GATACCGTAT
 2951 CCGGAGCGGC CGCCGCGTTA GCCGAGCGCC CCGCAGAGGA GTGGCTGGCG
25 3001 CGACCCCTGC CCGAGGGACT GCAGGCGITC GGGGCCGTCC TCGTAGACGC
 3051 CCATCGGCGC ATCACCGACC CGGAGAGGGA CATCCAGGAC TTTGTTCTCA
 3101 CCGCCGAACG GAGCAGACAC CCGCGCGCGT ACACCAACAA GCGCCTGGCC
30 3151 CACCTGACGG TGTATTACAA GCTCATGGCC CGCCGCGCGC AGGTCCCGTC
 3201 CATCAAGGAC CGGATCCCGT ACGTGATCGT GGCCCAGACC CGCGAGGTAG
35 3251 AGGAGACGGT CGCGCGGCTG GCCGCCCTCC GCGAGCTAGA CGCCGCCGCC
 3301 CCAGGGGACG AGCCCGCCCC CCCCAGGGCC CTGCCCTCCC CGGCCAAGCG
 3351 CCCCCGGGAG ACGCCGTCGC CTGCCGACCC CCCGGGAGGC GCGTCCAAGC
40 3401 CCCGCAAGCT GCTGGTGTCC GAGCTGGCCG AGGATCCCGC ATACGCCATT
 3451 GCCCACGGCG TCGCCCTGAA CACGGACTAT TACTTCTCCC ACCTGTTGGG
45 3501 GGCGGCGTGC GTGACATTCA AGGCCCTGTT TGGGAATAAC GCCAAGATCA
 3551 CCGAGAGTCT GTAAAAAAGG TTTATTCCCG AAGTGTGGCA CCCCCCGGAC
 3601 GACGTGGCCG CGCGGCTCCG GACCGCAGGG TTCGGGGCGG TGGGTGCCGG
50 3651 CGCTACGGCG GAGGAAACTC GTCGAATGTT GCATAGAGCC TTTGATACTC
 3701 TAGCATGA

5 **Figure 5T DNA and amino acid sequence list**

SEQ.ID.NO. 10 Amino acid sequence of DNA polymerase for HSV1-DJL-M1

1	MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGRGPPLC RQNFYNPYLA
10	51 PVGTQQKPTG PTQRHTYYSE CDEFRIFIAPR VLDEDAPPEK RAGVHDGHLK
	101 RAPKVYCGGD ERDVLRVGSG GFWPRRSRLW GGVDHAPAGF NPTVTVFHVY
15	151 DILENVEHAY GMRAAQFHAR FMDAITPTGT VITLLGLTPE GHRVAVHVG
	201 TRQYFYMNKE EVDRHLQCRA PRDLCEMAA ALRESPGASF RGISADHFEA
	251 EVVERTDVYY YETRPALFYR VYVRSGRVLS YLCDNFCPAI KKYEGGVDAT
20	301 TRFILDNPFG VTFGWYRLKP GRNNNTLAQPR APMAFGTSSD VEFNCTADNL
	351 AIEGGMSDLP AYKLMCFDIE CKAGGEDELA FPVAGHPEDL VIQISCLLYD
25	401 LSTTALEHVL LFSLGSCDLP ESHLNELAAR GLPTPVVLEF DSEFEMILLAF
	451 MTLVKQYGPE FVTGYNIINF DWPFLLAKLT DIYKVPLDGY GRMNGRGVFR
	501 VWDIGQSHFQ KR SKIKVNGM VNIDMYGIIT DKIKLSSYKL NAVA EAVLKD
30	551 KKKDLSYRDI PTYYAAGPAQ RG VIGEYCIQ DSLLVGQLFF KFLPHLELSA
	601 VARLAGINIT RTIYDGQQIR VFTCLLRLAD QKG FILPDTQ GRFRGAGGEA
35	651 PKRPAAARED EERPEEEGED ENEREEGGE REPEGARETA GRHVGYQGAR
	701 VLDPTSGFHV NPVVVFDFAS LYP SIIQAHN LCFSTLSLRA DAVA HLEAGK
	751 DYLEIEVGGR RLFFVKAHVR ESLLSILLRD WLAMRKQIRS RIPQSSPEEA
40	801 VLLDKQQAAI KVVCNSVYGF TGAQHGLLPC LHVAATVTTI GREMLLATRE
	851 YVHARWAAFE QLLADFPPEAA DMRAPGPYSM RIIYGDTDSI FVLCRGLTAA
	901 GLTAVGDKMA SHISRALFLP PIKLECEKTF TKLLLIAKKK YIGVIYGGKM
45	951 LIKGVDLVRK NNCAFINRTS RALVDLLFYD DT VSGAAAAL AERPAEEWLA
	1001 RPLPEGLQAF GAVLVDAHRR ITDPERDIQD FVLTAELSRH PRAYTNKRLA
50	1051 HLT VYYKLMA RRAQVPSIKD RIPYVIVAQT REVEETVARL AALRELDAAA
	1101 PGDEPAPPAA LPSPAKRPRE TPSPADPPGG ASKPRKLLVS ELAEDPAYAI
	1151 AHGVALNTDY YFSHLLGAAC VTFKALFGNN AKITESLLKR FIPEVWHPPD
55	1201 DVAARLRTAG FGAVGAGATA EETRRMLHRA FDTLA*

5 **Figure 5U DNA and amino acid sequence list**

SEQ.ID.NO. 11 DNA sequence of DNA polymerase gene for HMCV-AD169-M1

10 1 ATGTTTTCA ACCCGTATCT GAGCGGCCGG GTGACCGGCG GTGCGGTCGC
 51 GGGTGGCCGG CGTCAGCGTT CGCAGCCGG CTCCGCGCAG GGCTCGGGCA
 101 AGCGGCCGCC ACAGAAACAG TTTTGCAGA TCGTGCCGCG AGGTGTCATG
 15 151 TTGACGGTC AGACGGGGTT GATCAAGCAT AAGACGGGAC GGCTGCCTCT
 201 CATGTTCTAT CGAGAGATTAA AACATTTGTT GAGTCATGAC ATGGTTTGGC
 20 251 CGTGTCCCTTG GCGCGAGACC CTGGTGGTC GCGTGGTGGG ACCTATTCTG
 301 TTTCACACCT ACGATCAGAC GGACGCCGTG CTCTTCTTCG ACTCGCCGA
 351 AAACGTGTCG CCGCGCTATC GTCAGCATCT GGTGCCTTCG GGGAACGTGT
 25 401 TGCGTTCTT CGGGGCCACA GAACACGGCT ACAGTATCTG CGTCAACGTT
 451 TTCGGGCAGC GCAGCTACTT TTACTGTGAG TACAGCGACA CCGATAGGCT
 30 501 GCGTGAGGTC ATTGCCAGCG TGGGCGAACT AGTGCCGAA CCGCGGACGC
 551 CATA CGCCGT GTCTGTCACCG CCGGCCACCA AGACCTCCAT CTATGGGTAC
 601 GGGACGCGAC CCGTGCCCGA TTTGCAGTGT GTGTCTATCA GCAACTGGAC
 35 651 CATGCCAGA AAAATCGGCG AGTATCTGCT GGAGCAGGGT TTTCCCGTGT
 701 ACGAGGTCCG TGTGGATCCG CTGACCGT TGGTCATCGA TCGCGGATC
 40 751 ACCACGTTCG GCTGGTGCTC CGTGAATCGT TACGACTGGC GGCAGCAGGG
 801 TCGCGCGTCG ACTTGTGATA TCGAGGTAGA CTGCGATGTC TCTGACCTGG
 851 TGGCTGTGCC CGACGACAGC TCGTGGCCGC GCTATCGATG CCTGTCCTTC
 45 901 GATATCGAGT GCATGAGCGG CGAGGGTGGT TTTCCCTGCG CCGAGAAGTC
 951 CGATGACATT GTCATTCAAGA TCTCGTGCCT GTGCTACGAG ACGGGGGGAA
 50 1001 ACACCGCCGT GGATCAGGGG ATCCCAAACG GGAACGATGG TCGGGGCTGC
 1051 ACTTCGGAGG GTGTGATCTT TGGGCACTCG GGTCTTCATC TCTTACGAT
 1101 CGGCACCTGC GGGCAGGTGG GCCCAGACGT GGACGTCTAC GAGTTCCCTT
 55 1151 CCGAATACGA GCTGCTGCTG GGCTTATGC TTTTCTTCAC ACGGTACGCG
 1201 CCGGCCTTG TGACCGGTTA CAACATCAAC TCITTTGACT TGAAGTACAT

60

5 **Figure 5V DNA and amino acid sequence list**

1251 CCTCACGCGT CTCGAGTACC TGTATAAGGT GGACTCGCAG CGCTTCTGCA
10 1301 AGTTGCCTAC GGCGCAGGGC GGCGTITCT TTTACACAG CCCCGCCGTG
1351 GGTTTTAAGC GGCAGTACGC CGCOGCTTT CCCTCGGCTT CTCACAACAA
15 1401 TCCGGCCAGC ACGGCCGCCA CCAAGGTGTA TATTGCGGGT TCGGTGGTTA
1451 TCGACATGTA CCCTGTATGC ATGGCCAAGA CTAACTCGCC CAACTATAAG
20 1501 CTCAACACTA TGGCCGAGCT TTACCTGOGG CAACGCAAGG ATGACCTGTC
1551 TTACAAGGAC ATCCCGCGTT GTTCTGTGGC TAATGCCAGG GGCCGCGCCC
1601 AGGTAGGCCG TTACTGTCTG CAGGACGCCG TATTGGTGCG CGATCTGTT
25 1651 AACACCATTA ATTTCACTA CGAGGCCGGG GCCATCGCGC GGCTGGCTAA
1701 AATTCCGTTG CGCGTGTCA TCTTGACGG ACAGCAGATC CGTATCTACA
1751 CCTCGCTGCT GGACGAGTGC GCCTGCCGCG ATTATCCT GCCCAACCAC
30 1801 TACAGCAAAG GTACGACGGT GCCCGAAACG AATAGCGTTG CTGTGTCACC
1851 TAACGCTGCT ATCATCTCTA CCGCCGCTGT GCCCGGCGAC GCGGGTTCTG
35 1901 TGGCGGCTAT GTTCAGATG TCGCCGCCCT TGCAATCTGC GCCGTCCAGT
1951 CAGGACGGCG TTTCACCCGG CTCCGGCAGT AACAGTAGTA GCAGCGTCGG
2001 CGTTTCAGC GTCGGCTCCG GCAGTAGTGG CGCGTCGGC GTTCCAACG
40 2051 ACAATCACGG CGCCGGCGGT ACTGCCGGGG TTTCGTACCA GGGGCCACG
2101 GTGTTGAGC CCGAGGTGGG TTACTACAAC GACCCCGTGG CCGTGTTCGA
45 2151 CTTGCCAGC CTCTACCCCT CCATCATCAT GGCCCACAAC CTCTGCTACT
2201 CCACCCCTGCT GGTGCCGGGT GGCGAGTACC CTGTGGACCC CGCCGACGTA
2251 TACAGCGTCA CGCTAGAGAA CGGCGTGACC CACCGCTTG TGCGTGCTTC
50 2301 GGTGCGCGTC TCGGTGCTCT CGGAACGTGCT CAACAAGTGG GTTTCGCA
2351 GGCCTGCCGT GCGCGAATGC ATGCGCGAGT GTCAAGACCC TGTGCGCCGT
2401 ATGCTGCTCG ACAAGGAACA GATGGCGCTC AAAGTAACGT GCAACGCTTT
55 2451 CTACGGTTTT ACCGGCGCGC TGAACGGTAT GATGCCGTGT CTGCCCATCG
2501 CCGCCAGCAT CACGCGCATC GGTGCGACG TGCTAGAGCG CACGGCGCGG

Figure 5W DNA and amino acid sequence list

10 2551 TTCATCAAAG ACAACTTTTC AGAGCCGTGT TTTTGACAA ATTTTTTAA
 2601 TCAGGAAGAC TATGTAGTGG GAACGCGGGA GGGGGATTG GAGGAGAGCA
 2651 GCGCGTTACC GGAGGGGCTC GAAACATCGT CAGGGGGCTC GAACGAACGG
 15 2701 CGGGTGGAGG CGCGGGTCAT CTACGGGGAC ACGGACAGCG TGTTGTCCG
 2751 CTTTGTGGC CTGACGCCGC AGGCTCTGGT GGCGCGTGGG CCCAGCCTGG
 20 2801 CGCACTACGT GACGGCCTGT CTTTTGTGG AGCCCGTCAA GCTGGAGTTT
 2851 GAAAAGGTCT TCGTCTCTCT TATGATGATC TGCAAGAAC GTTACATCGG
 25 2901 CAAAGTGGAG GGCGCCTCGG GTCTGAGCAT GAAGGGCGTG GATCTGGTGC
 2951 GCAAGACGGC CTGCGAGTTTC GTCAAGGGCG TCACGCGTGA CGTCCTCTCG
 3001 CTGCTCTTG AGGATCGCGA GGTCTCGGAA GCAGCCGTGC GCCTGTGCG
 30 3051 CCTCTCACTC GATGAAGTCA AGAAGTACGG CGTGCCACGC GGTTTCTGGC
 3101 GTATCTTACG CCGCTTGGTG CAGGCCCGCG ACGATCTGTA CCTGCACCGT
 35 3151 GTGCGTGTGCG AGGACCTGGT GCTTCGTGCG GTGCTCTCTA AGGACATCTC
 3201 GCTGTACCGT CAATCTAACCC TGCCGCACAT TGCCGTCAATT AAGCGATTGG
 3251 CGGCCCGTTC TGAGGAGCTA CCCTCGTGC GGGATCGGGT CTTTACGTT
 40 3301 CTGACGGCGC CCGGTGTCCG GACGGCGCCG CAGGGTTCCCT CCGACAACGG
 3351 TGATTCTGTA ACCGCCGGCG TGTTTCCCG GTCGGACGCG ATTGATGGCA
 45 3401 CGGACGACGA CGCTGACGGC GGCGGGTAG AGGAGAGCAA CAGGAGAGGA
 3451 GGAGAGCCGG CAAAGAAGAG GGCGCGGAAA CCACCGTCGG CCGTGTGCAA
 3501 CTACGAGGTA GCCGAAGATC CGAGCTACGT GCGCGAGCAC GGC GTGCCA
 50 3551 TTCACGCCGA CAAGTACTTT GAGCAGGTTTC TCAAGGCTGT AACTAACGTG
 3601 CTGTCGCCCG TCTTTCCCGG CGCGAAACC GCGCGCAAGG ACAAGTTTT
 3651 GCACATGGTG CTGCCGCCGC GCTTGCACCTT GGAGCCGGCT TTTCTGCCGT
 55 3701 ACAGTGTCAA GGCGCACGAA TGCTGTTGA

5 **Figure 5X DNA and amino acid sequence list****SEQ.ID.NO.12 Amino acid sequence of DNA polymerase f r HCMV-AD169-M1**

10 1 MFFNPYLSGG VTGGAVAGGR RQRSQPGSAQ GSGKRPPQKQ FLQIVPRGVM
 51 FDGQTGLIKH KTGRLPLMFY REIKHLLSHD MVWPCPWRET LVGRVVGPIR
 101 FHTYDQTDAV LFFDSPENVS PRYRQHLVPS GNVLRFFGAT EHGYSICVNV
 15 151 FGQR SYFYCE YSDTDR LREV IASVGELVPE PRTPYAVSVT PATKTSIYGY
 201 GTRPV PDLQC VSISNWTMAR KIGEYLLEQG FPVYEV RVDP LTRLVIDRRI
 20 251 TTFGWC SVNR YDWRQQGRAS TCDIEVDCDV SDLVA VPDDS SWPRYRCLSF
 301 DIECMSGEGG FPCA EKSDDI VIQISCVCYE TGGNTAVDQG IPNGNDGRGC
 351 TSEGVI FGHS GLHLFTIGTC GQVGPDVDVY EFPSEYELL GFMLFFQRYA
 25 401 PAFVTGYNIN SFDLKYILTR LEYLYKVDSQ RFCKLPTA QG GRFFLHSPAV
 451 GFKRQYAAAF PSASHNNPAS TAATK VYIAG SVVIDMYPVC MAKTN SPNYK
 30 501 LNTMAELYLR QRKDDLSYKD IPRCFVANAE GRAQVGRYCL QDAVLVRDLF
 551 NTINFHYEAG AIARLAKIPL RRVIFDGQQI RIYTSLLDEC ACRDFILPNH
 601 YSKGTTVPET NSVAVSPNAA IISTAAVPGD AGSVAAMFQM SPPLQSAPSS
 35 651 QDGVSPGS GS NSSSSVG VFS VGSGSSGGVG VSNDNHGAGG TAAVSYQGAT
 701 VFEPEVGYYN DPVA VFDFAS L YPSII MAHN LCYST LLVPG GEYPVDPADV
 40 751 YSVTLENGVT HRFVRAS VRV SVLSELLNKW VSQRRAVREC MRECQDPVRR
 801 MLLDKEQMAL KVTCNAFYGF TGALNGMMPC LPIAASITRI GRDMLERTAR
 851 FIKDNFSEPC FLHNNFNQED YVVG TREGDS EESSALPEG ETSSGGSN ER
 45 901 RVEARV IYGD TDSVFVRFRG LTPQALVARG PSLAHYVTAC LFVEPVKLEF
 951 EKVFVSLMMI CKKRYIGKVE GASGLSMKG V DLVRKTACEF VKGVTRDVLS
 50 1001 LLFEDREVSE AAVRLSRLSL DEVKKYGVPR GFWRILRRLV QARDDLYLHR
 1051 VRVEDLV LSS VLSKDISLYR QSNLPHIAVI KRLAARSEEL PSVGDRV FYV
 1101 LTAPGVRTAP QGSSDNGDSV TAGVVRSDA IDGT DDDADG GGVEESNRRG
 55 1151 GEPACKRARK PPSAVCN YEV AEDPSYVREH GPVHADKYF EQVLKAVTNV
 1201 LSPVFP GGET ARKDKFLHMV LP RRLHLEPA FLPYSVKAHE CC*